GIT RE-BASIN: MERGING MODELS MODULO PERMUTATION SYMMETRIES

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ABSTRACT

The success of deep learning is due in large part to our ability to solve certain massive non-convex optimization problems with relative ease. Though non-convex optimization is NP-hard, simple algorithms - often variants of stochastic gradient descent - exhibit surprising effectiveness in fitting large neural networks in practice. We argue that neural network loss landscapes contain (nearly) a single basin after accounting for all possible permutation symmetries of hidden units a la Entezari et al. (2021). We introduce three algorithms to permute the units of one model to bring them into alignment with a reference model in order to merge the two models in weight space. This transformation produces a functionally equivalent set of weights that lie in an approximately convex basin near the reference model. Experimentally, we demonstrate the single basin phenomenon across a variety of model architectures and datasets, including the first (to our knowledge) demonstration of zero-barrier linear mode connectivity between independently trained ResNet models on CIFAR-10 and CIFAR-100. Additionally, we identify intriguing phenomena relating model width and training time to mode connectivity. Finally, we discuss shortcomings of the linear mode connectivity hypothesis, including a counterexample to the single basin theory.

1 Introduction

We investigate the unreasonable effectiveness of stochastic gradient descent (SGD) algorithms on the high-dimensional non-convex optimization problems of deep learning. In particular,

- 1. Why does SGD thrive in optimizing high-dimensional non-convex deep learning loss land-scapes despite being noticeably less robust in other non-convex optimization settings, like policy learning (Ainsworth et al., 2021), trajectory optimization (Kelly, 2017), and recommender systems (Kang et al., 2016)?
- 2. What are all the local minima? When linearly interpolating between initialization and final trained weights, why does the loss smoothly and monotonically decrease (Goodfellow & Vinyals, 2015; Frankle, 2020; Lucas et al., 2021; Vlaar & Frankle, 2021)?
- 3. How can two independently trained models with different random initializations and data batch orders inevitably achieve nearly identical performance? Furthermore, why do their training loss curves look identical?

We posit that the final phenomenon referenced in item 3 points to the existence of some yet uncharacterized invariance(s) in the training dynamics that cause independent training runs to exhibit almost identical characteristics. Hecht-Nielsen (1990) noted the permutation symmetries of hidden units in neural networks; briefly, one can swap any two units of a hidden layer in a network and – assuming weights are adjusted accordingly – network functionality will not change. Recently, Entezari et al. (2021) conjectured that these permutation symmetries may let us linearly connect points in weight space with no detriment to the loss.

Conjecture 1 (Permutation invariance, informal (Entezari et al., 2021)). Most SGD solutions belong to a set whose elements can be permuted so that no barrier (as in Definition 2.2) exists on the linear interpolation between any two permuted elements.

| ARCHITECTURE | NUM. PERMUTATION SYMMETRIES |
|----------------------------------|-----------------------------|
| MLP (3 layers, 512 width) | $10 \wedge 3498$ |
| VGG16 | $10 \land 35160$ |
| ResNet50 | $10 \wedge 55109$ |
| Atoms in the observable universe | 10 ∧ 82 |

Table 1: Permutation symmetries of deep learning models vs. an upper estimate on the number of atoms in the known, observable universe. Deep learning loss landscapes contain incomprehensible amounts of geometric repetition.

We refer to such solutions as being *linearly mode connected* (LMC) (Frankle et al., 2020). If true, Conjecture 1 will both materially expand our understanding of how SGD works in the context of deep learning and offer a credible explanation for the preceding phenomena, in particular.

Contributions. In this paper, we attempt to uncover what invariances may be responsible for the phenomena cited above and the unreasonable effectiveness of SGD in deep learning. We make the following contributions:

- 1. **Matching methods.** We propose three new algorithms, grounded in concepts and techniques from combinatorial optimization, to align the weights of two independently trained models. Where appropriate, we prove hardness results for these problems and propose approximation algorithms. Our fastest method identifies permutations in mere seconds on current hardware.
- 2. **Relationship to SGD.** We demonstrate by means of counterexample that linear mode connectivity is an emergent property of SGD training, not of model architectures. We connect this result to prior work on the implicit biases of SGD.
- 3. Experiments, including zero-barrier LMC for ResNets. Empirically, we explore the existence of linear mode connectivity modulo permutation symmetries in experiments across MLPs, CNNs, and ResNets trained on MNIST, CIFAR-10, and CIFAR-100. We contribute the first-ever demonstration of zero-barrier LMC between two independently trained ResNets. We explore the relationship between LMC and model width as well as training time. Finally, we show evidence of our methods' ability to combine models trained on independent datasets into a merged model that outperforms both input models in terms of test loss (but not accuracy) and is no more expensive in compute or memory than either input model.

2 BACKGROUND

Although our methods can be applied to arbitrary model architectures, we proceed with the multi-layer perceptron (MLP) for its ease of presentation (Bishop, 2007). Consider an *L*-layer MLP,

$$f(\boldsymbol{x};\Theta) = \boldsymbol{z}_{L+1}, \quad \boldsymbol{z}_{\ell+1} = \sigma(\boldsymbol{W}_{\ell}\boldsymbol{z}_{\ell} + \boldsymbol{b}_{\ell}), \quad \boldsymbol{z}_{1} = \boldsymbol{x},$$

where σ denotes an element-wise nonlinear activation function. Furthermore, consider a loss, $\mathcal{L}(\Theta)$, that measures the suitability of a particular set of weights Θ towards some goal, e.g., fitting to a training dataset.

Central to our investigation is the phenomenon of *permutation symmetries* of weight space. Given Θ , we can apply some permutation to the output features of any intermediate layer, ℓ , of the model, denoted by a permutation matrix $P \in S_d$, ¹

$$\boldsymbol{z}_{\ell+1} = \boldsymbol{P}^{\top} \boldsymbol{P} \boldsymbol{z}_{\ell+1} = \boldsymbol{P}^{\top} \boldsymbol{P} \sigma (\boldsymbol{W}_{\ell} \boldsymbol{z}_{\ell} + \boldsymbol{b}_{\ell}) = \boldsymbol{P}^{\top} \sigma (\boldsymbol{P} \boldsymbol{W}_{\ell} \boldsymbol{z}_{\ell} + \boldsymbol{P} \boldsymbol{b}_{\ell})$$

for σ , an element-wise operator. It follows that as long as we reorder the input weights of layer $\ell+1$ according to \boldsymbol{P}^{\top} , we will have a functionally equivalent model. To be precise, if we define Θ' to be identical to Θ with the exception of

$$\boldsymbol{W}_{\ell}' = \boldsymbol{P} \boldsymbol{W}_{\ell}, \quad \boldsymbol{b}_{\ell}' = \boldsymbol{P} \boldsymbol{b}_{\ell}, \quad \boldsymbol{W}_{\ell+1}' = \boldsymbol{W}_{\ell+1} \boldsymbol{P}^{\top},$$

¹We denote the set of all $d \times d$ permutation matrices – isomorphic to the symmetric group – as S_d , to the possible chagrin of pure mathematicians.

then the two models are functionally equivalent: $f(x; \Theta) = f(x; \Theta')$ for all inputs x. This implies that for any trained weights Θ , there is an entire equivalence class of functionally equivalent weight assignments, not just one such Θ , and convergence to any one specific element of this equivalence class, as opposed to any others, is determined only by random seed. We denote a functionality-preserving permutation of weights as $\pi(\Theta)$.

Consider the task of reconciling the weights of two, independently trained models, A and B, with weights Θ_A and Θ_B , respectively, such that we can linearly interpolate between them. We assume that models A and B were trained with equivalent architectures but different random initializations, data orders, and potentially different hyperparameters or datasets, as well. Our central question is: Given Θ_A and Θ_B , can we identify some π such that when linearly interpolating between Θ_A and $\pi(\Theta_B)$, all intermediate models enjoy performance similar to Θ_A and Θ_B ?

We base any claims of loss landscape convexity on the usual definition of multi-dimensional convexity in terms of one-dimensional convexity per

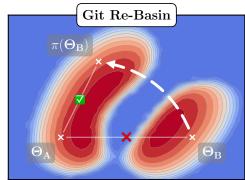


Figure 1: Git Re-Basin merges models by teleporting solutions into a single basin. Θ_B is permuted into $\pi(\Theta_B)$ so that it lies in the same basin as Θ_A .

Definition 2.1 (Convexity). A function $f: \mathbb{R}^D \to \mathbb{R}$ is convex if every one-dimensional slice is convex, i.e., for all $x, y \in \mathbb{R}^D$, the function $g(\lambda) = f((1 - \lambda)x + \lambda y)$ is convex in λ .

Due to Definition 2.1, it suffices to show that arbitrary one-dimensional slices of a function are convex in order to reason about the convexity of complex, high-dimensional functions. In practice, we rarely observe perfect convexity but instead hope to approximate it as closely as possible. Following Draxler et al. (2018), Entezari et al. (2021), Frankle et al. (2020) and others, we measure approximations to convexity via "barriers."

Definition 2.2 (Loss barrier (Frankle et al., 2020)). Given two points Θ_A , Θ_B such that $\mathcal{L}(\Theta_A) \approx \mathcal{L}(\Theta_B)$, the *loss barrier* is defined as $\max_{\lambda \in [0,1]} \mathcal{L}((1-\lambda)\Theta_A + \lambda\Theta_B) - \frac{1}{2}(\mathcal{L}(\Theta_A) + \mathcal{L}(\Theta_B))$.

Loss barriers are non-negative, with zero indicating an interpolation of flat or positive curvature.

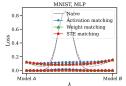
3 Permutation Selection Methods

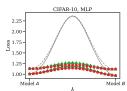
We introduce three methods of matching units between model A and model B. Further, we present an extension to simultaneously merging multiple models in Appendix A.6 and an appealing but failed method in Appendix A.7.

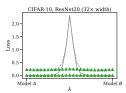
3.1 MATCHING ACTIVATIONS

Following the classic Hebbian mantra, "[neural network units] that fire together, wire together" (Hebb, 2005), we propose associating units across two models by performing regression between their activations. Matching activations between models is compelling since it captures the intuitive notion that two models must learn similar features to accomplish the same task (Li et al., 2016). Provided activations for each model, we aim to associate each unit in A with a unit in B. It stands to reason that a linear relationship may exist between the activations of the two models. We fit this into the regression framework by constraining ordinary least squares (OLS) to solutions in the set of permutation matrices, S_d . For activations of the ℓ 'th layer, let $\mathbf{Z}^{(A)}$, $\mathbf{Z}^{(B)} \in \mathbb{R}^{d \times n}$ denote the d-dim. activations for all n training data points in models A and B, respectively. Then,

$$P_{\ell} = \underset{P \in S_d}{\operatorname{arg \, min}} \sum_{i=1}^{n} \| Z_{:,i}^{(A)} - P Z_{:,i}^{(B)} \|^{2} = \underset{P \in S_d}{\operatorname{arg \, max}} \langle P, Z^{(A)} (Z^{(B)})^{\top} \rangle_{F}, \tag{1}$$







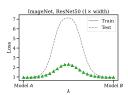


Figure 2: Linear mode connectivity is possible after permuting. Loss when interpolating between models trained on MNIST, CIFAR-10, and ImageNet. In all cases we can significantly improve over naïve interpolation. Straight-through estimator matching performs best but is very computationally expensive. Weight and activation matching perform similarly, although weight matching is orders of magnitude faster and does not rely on the input data distribution. We hypothesize that the ImageNet barrier could be reduced by increasing the model width as in Section 5.3.

where $\langle \boldsymbol{A}, \boldsymbol{B} \rangle_F = \sum_{i,j} A_{i,j} B_{i,j}$ denotes the Frobenius inner product between real-valued matrices \boldsymbol{A} and \boldsymbol{B} . Conveniently, eq. (1) constitutes a "linear assignment problem" (LAP) (Bertsekas, 1998) for which efficient, practical algorithms are known. Having solved this assignment problem on each layer, we can then permute the weights of model \boldsymbol{B} to match model \boldsymbol{A} as closely as possible

$$oldsymbol{W}_\ell' = oldsymbol{P}_\ell oldsymbol{W}_\ell^{(B)} oldsymbol{P}_{\ell-1}^ op, \quad oldsymbol{b}_\ell' = oldsymbol{P}_\ell oldsymbol{b}_\ell^{(B)}$$

for each layer ℓ , producing weights Θ' with activations that align as closely possible with Θ_A .

Computationally, this entire process is relatively lightweight: the $\mathbf{Z}^{(A)}$ and $\mathbf{Z}^{(B)}$ matrices can be computed in a single pass over the training dataset, and, in practice, a full run through the training dataset may be unnecessary. Solving eq. (1) is possible due to well-established, polynomial-time algorithms for solving the linear assignment problem (Kuhn, 2010; Jonker & Volgenant, 1987; Crouse, 2016). Also, conveniently, the activation matching at each layer is independent of the matching at every other layer, resulting in a separable and straightforward optimization problem; this advantage will not be enjoyed by the following methods.

Dispensing with regression, one could similarly associate units by matching against a matrix of cross-correlation coefficients in place of $\mathbf{Z}^{(A)}(\mathbf{Z}^{(B)})^{\top}$. We observed correlation matching to work equally well but found OLS regression matching to be more principled and easier to implement.

3.2 MATCHING WEIGHTS

Instead of associating units by their activations, we could alternatively inspect the weights of the model itself. Consider the first layer weights, W_1 ; each row of W_1 corresponds to a single feature. If two such rows were equal, they would compute exactly the same feature (ignoring bias terms for the time being). And, if $[W_1^{(A)}]_{i,:} \approx [W_1^{(B)}]_{j,:}$, it stands to reason that units i and j should be associated. Extending this idea to every layer, we are inspired to pursue the optimization

$$\underset{\pi}{\arg\min} \ \| \operatorname{vec}(\Theta_A) - \operatorname{vec}(\pi(\Theta_B)) \|^2 = \underset{\pi}{\arg\max} \ \operatorname{vec}(\Theta_A) \cdot \operatorname{vec}(\pi(\Theta_B)).$$

We can re-express this in terms of the full weights,

$$\underset{\pi=\{P_L\}}{\operatorname{arg\,max}} \langle \boldsymbol{W}_1^{(A)}, \, \boldsymbol{P}_1 \boldsymbol{W}_1^{(B)} \rangle_F + \langle \boldsymbol{W}_2^{(A)}, \, \boldsymbol{P}_2 \boldsymbol{W}_2^{(B)} \boldsymbol{P}_1^\top \rangle_F + \dots + \langle \boldsymbol{W}_L^{(A)}, \, \boldsymbol{W}_L^{(B)} \boldsymbol{P}_{L-1}^\top \rangle_F, \quad (2)$$

resulting in another matching problem. We term this formulation the "sum of bilinear assignments problem" (SOBLAP). Unfortunately, this matching problem is thornier than the classic linear assignment matching problem presented in eq. (1). Unlike LAP, we are interested in permuting both the rows and columns of $\boldsymbol{W}_{\ell}^{(B)}$ to match $\boldsymbol{W}_{\ell}^{(A)}$, which fundamentally differs from permuting only rows or only columns. We formalize this difficulty as follows.

Lemma 1. The sum of a bilinear assignments problem (SOBLAP) is NP-hard and admits no polynomial-time constant-factor approximation scheme for L > 2.

Lemma 1 contrasts starkly with classical LAP, for which polynomial-time algorithms are known.

Undeterred, we propose a approximation algorithm for SOBLAP. Looking at a single P_{ℓ} while holding the others fixed, we observe that the problem can be reduced to a classic LAP,

$$\begin{aligned} \arg\max_{\boldsymbol{P_{\ell}}} & \langle \boldsymbol{W}_{\ell}^{(A)}, \boldsymbol{P_{\ell}} \boldsymbol{W}_{\ell}^{(B)} \boldsymbol{P}_{\ell-1}^{\top} \rangle_{F} + \langle \boldsymbol{W}_{\ell+1}^{(A)}, \boldsymbol{P_{\ell+1}} \boldsymbol{W}_{\ell+1}^{(B)} \boldsymbol{P}_{\ell}^{\top} \rangle_{F} \\ &= \arg\max_{\boldsymbol{P_{\ell}}} & \langle \boldsymbol{P_{\ell}}, \boldsymbol{W}_{\ell}^{(A)} \boldsymbol{P_{\ell-1}} (\boldsymbol{W}_{\ell}^{(B)})^{\top} + (\boldsymbol{W}_{\ell+1}^{(A)})^{\top} \boldsymbol{P_{\ell+1}} \boldsymbol{W}_{\ell+1}^{(B)} \rangle_{F}. \end{aligned}$$

This leads to a convenient coordinate descent algorithm: go through each layer and greedily select its best P_{ℓ} . Repeat until convergence. We present this in Algorithm 1.

Algorithm 1: PERMUTATION COORDINATE DESCENT

```
Given: Model weights \Theta_A = \left\{ \boldsymbol{W}_1^{(A)}, \dots, \boldsymbol{W}_L^{(A)} \right\} and \Theta_B = \left\{ \boldsymbol{W}_1^{(B)}, \dots, \boldsymbol{W}_L^{(B)} \right\}

Result: A permutation \pi = \left\{ \boldsymbol{P}_1, \dots, \boldsymbol{P}_{L-1} \right\} of \Theta_B such that \text{vec}(\Theta_A) \cdot \text{vec}(\pi(\Theta_B)) is approximately maximized.
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\begin{split} \textbf{Initialize:} & \ \boldsymbol{P}_1 \leftarrow \boldsymbol{I}, \dots, \boldsymbol{P}_{L-1} \leftarrow \boldsymbol{I} \\ \textbf{repeat} \\ & \ | \ \ \textbf{for} \ \ell \in \texttt{RANDOMPERMUTATION}(1, \dots, L-1) \ \textbf{do} \\ & \ | \ \ \boldsymbol{P}_\ell \leftarrow \texttt{SOLVELAP} \left( \boldsymbol{W}_\ell^{(A)} \boldsymbol{P}_{\ell-1} (\boldsymbol{W}_\ell^{(B)})^\top + (\boldsymbol{W}_{\ell+1}^{(A)})^\top \boldsymbol{P}_{\ell+1} \boldsymbol{W}_{\ell+1}^{(B)} \right) \\ & \ \ \textbf{end} \\ & \ \ \textbf{until} \ \textit{convergence} \end{split}
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Although we present Algorithm 1 in terms of an MLP without bias terms, in practice our implementation can handle the weights of models of arbitrary architectures, including bias terms, residual connections, convolutional layers, attention mechanisms, and so forth. We propose an extension of Algorithm 1 to merging more than two models at a time in Appendix A.6.

Lemma 2. Algorithm 1 terminates.

Our experiments showed this algorithm to be fast in terms of both iterations necessary for convergence and wall-clock time, generally on the order of seconds to a few minutes.

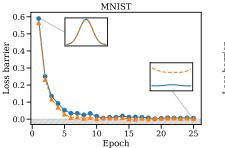
Unlike the activation matching method presented in Section 3.1, weight matching ignores the data distribution entirely. Ignoring the input data distribution and therefore the loss landscape handicaps weight matching but allows it to be much faster. We therefore anticipate its potential application in fields such as finetuning (Devlin et al., 2019; Wortsman et al., 2022b;a), federated learning (McMahan et al., 2017; Konečný et al., 2016a;b), and model patching (Matena & Raffel, 2021; Sung et al., 2021; Raffel, 2021). In practice, we found weight matching to be surprisingly competitive with data-aware methods. Section 5 studies this trade-off.

3.3 LEARNING PERMUTATIONS WITH A STRAIGHT-THROUGH ESTIMATOR

Inspired by the success of straight-through estimators (STEs) in other discrete optimization problems (Bengio et al., 2013; Kusupati et al., 2021; Rastegari et al., 2016; Courbariaux & Bengio, 2016), we attempt here to "learn" the ideal permutation of weights $\pi(\Theta_B)$. Specifically, our goal is to optimize

$$\min_{\tilde{\Theta}_B} \mathcal{L}\left(\frac{1}{2}\left(\Theta_A + \operatorname{proj}\left(\tilde{\Theta}_B\right)\right)\right), \quad \operatorname{proj}(\Theta) \triangleq \arg\max_{\pi} \operatorname{vec}(\Theta) \cdot \operatorname{vec}(\pi(\Theta_B)), \quad (3)$$

where $\tilde{\Theta}_B$ denotes an approximation of $\pi(\Theta_B)$, allowing us to implicitly optimize π . However, eq. (3) involves inconvenient non-differentiable projection operations, $\operatorname{proj}(\cdot)$, complicating the optimization. We overcome this via a "straight-through" estimator: we parameterize the problem in terms of a set of weights $\tilde{\Theta}_B \approx \pi(\Theta_B)$. In the forward pass, we project $\tilde{\Theta}_B$ to the closest realizable $\pi(\Theta_B)$. In the backwards pass, we then switch back to the unrestricted weights $\tilde{\Theta}_B$. In this way, we



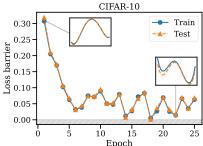


Figure 3: Linear mode connectivity is challenging at initialization. We show loss barriers per training time for MLPs trained on MNIST (left) and CIFAR-10 (right). Loss interpolation plots are inlaid to highlight results in initial and later epochs. LMC manifests gradually throughout training. We hypothesize that the variance in CIFAR-10 training is higher due to our MLP architecture being under-powered relative to the dataset. (Y-axis scales differ in each inlaid plot.)

are guaranteed to stay true to the projection constraints in evaluating the loss but can still compute usable gradients at our current parameters, $\tilde{\Theta}_B$.²

Conveniently, we can re-purpose Algorithm 1 to solve $\operatorname{proj}(\tilde{\Theta}_B)$. Furthermore, we found that initializing $\tilde{\Theta}_B = \Theta_A$ performed better than random initialization. This is to be expected immediately at initialization since the initial matching will be equivalent to the weight matching method of Section 3.1. However, it is not immediately clear why these solutions continue to outperform a random initialization asymptotically.

Unlike the aforementioned methods, Algorithm 2 attempts to explicitly "learn" the best permutation π using a conventional training loop. By initializing to the weight matching solution of Section 3.2 and leveraging the data distribution as in Section 3.1, it seeks to offer a best-of-both-worlds solution. However, this comes at a very steep computational cost relative to the other two methods.

4 Interlude: A Counterexample

Before arguing for the presence of linear mode connectivity, we step back to consider its limitations. In Appendix A.3 we present a counterexample proving that there exist models that do not enjoy LMC under any permutation of weights. Since our two networks have small width, we can simply inspect all four possible permutations of the intermediate units. We visualize this in Figure 6. Critically, no permutation results in LMC.

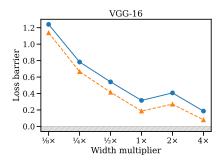
We present this counterexample in part to establish some basic intuition for Conjecture 1 but more importantly to highlight that any success interpolating between permuted networks is due to inherent bias in the optimization algorithms used, not the network architectures themselves. To the extent we can achieve LMC in practice, it follows that it is an artifact of our optimization methods, not our architectures. SGD is implicitly biased towards solutions admitting LMC.

We also note that there are invariances beyond permutation symmetries: It is possible to move features between layers, re-scale layers, and so forth. Prior works noted the feature/layer association (Nguyen et al., 2021) and re-scaling invariances (Ainsworth et al., 2018), albeit in different contexts. The importance of these other symmetries and their interplay with SGD remains unclear.

5 EXPERIMENTS

Our base methodology is to separately train two models, A and B, starting from different random initializations and with different random batch orders, resulting in trained weights Θ_A and Θ_B ,

²Note again that projecting according to inner product distance is equivalent to projecting according to the L_2 distance when parameterizing the estimator based on the B endpoint. We also experimented with learning the midpoint directly, $\tilde{\Theta} \approx \frac{1}{2}(\Theta_A + \pi(\Theta_B))$, in which case the L_2 and inner product projections diverge. In testing all possible variations, we found that optimizing the B endpoint had a slight advantage, but all possible variations performed admirably.



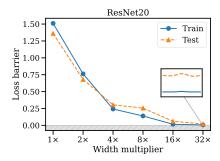


Figure 4: **Wider models exhibit better linear mode connectivity.** Training convolutional and ResNet architectures on CIFAR-10, we ablate their width and visualize loss barriers after weight matching. Notably, we achieve zero-barrier linear mode connectivity between ResNet models, the first such demonstration to the authors' knowledge.

respectively. We then evaluate slices through the loss landscape, $\mathcal{L}((1-\lambda)\Theta_A + \lambda\pi(\Theta_B))$ for $\lambda \in [0,1]$, where π is selected according to the methods presented in Section 3.³ Ideally, we seek a completely flat or even convex one-dimensional slice. As discussed in Section 2, the ability to exhibit this behavior for arbitrary Θ_A, Θ_B empirically suggests that the loss landscape contains only a single basin modulo permutation symmetries.

We remark that a failure to find a π such that linear mode connectivity holds cannot rule out the existence of a satisfactory permutation. Given the astronomical number of permutation symmetries, Conjecture 1 is essentially impossible to disprove for any realistically wide model architecture.

5.1 Loss Landscapes Before and After Matching

We present results for models trained on MNIST (LeCun et al., 1998), CIFAR-10 (Krizhevsky, 2009), and ImageNet (Deng et al., 2009) in Figure 2. Naïve interpolation $(\pi(\Theta_B) = \Theta_B)$ substantially degrades performance when interpolating. On the other hand, the methods introduced in Section 3 can achieve much better barriers. We achieve zero-barrier linear mode connectivity on MNIST with all three methods, although activation matching performs just slightly less favorably than weight matching and straight-through estimator (STE) matching. We especially note that the test loss landscape becomes convex after applying our weight matching and STE permutations! In other words, our interpolation actually yields a merged model that outperforms both models A and B. We elaborate on this phenomenon in Section 5.4 and Appendix A.6.

On ImageNet we fall short of zero-barrier connections, although we do see a 67% decrease in barrier relative to naïve interpolation. As we demonstrate in Section 5.3, we can achieve zero-barrier LMC on CIFAR-10 with large ResNet models. Therefore, we hypothesize that the presence of LMC depends on the model having sufficient capacity (esp. width) to capture the complexity of the input data distribution, and that ImageNet results could be improved by expanding model width.

STE matching, the most expensive method, produces the best solutions. Somewhat surprising, however, is that the gap between STE and the other two methods is relatively small. In particular, it is remarkable how well Algorithm 1 performs without access to the input data at all. We found that weight matching offered a compelling balance between computational cost and performance: It runs in mere seconds (on modern hardware) and produces high-quality solutions.

5.2 Onset of Mode Connectivity

Given the results of Section 5.1, it may be tempting to conclude that the entirety of weight space contains only a single basin modulo permutation symmetries. However, we found that linear mode connectivity is an emergent property of SGD training, and we were unable to uncover it early in training. We explore the emergence of LMC in Figure 3. Concurrent to our work, Benzing et al. (2022) showed that LMC at initialization is possible using a permutation found at the end of training.

³We also experimented with spherical linear interpolation ("slerp") and found it to perform slightly better than linear interpolation; however, the difference was not sufficiently significant to warrant diverging from the pre-existing literature.

Note that the final inlaid interpolation plot in Figure 3(right) demonstrates an important shortcoming of the loss barrier metric, i.e., the interpolation includes points with lower loss than either of the two models. However, the loss barrier is still positive due to non-negativity, as mentioned in Section 2.

5.3 EFFECT OF MODEL WIDTH

Conventional wisdom maintains that wider architectures are easier to optimize (Jacot et al., 2018; Lee et al., 2019). We now investigate whether they are also easier to linearly mode connect. We train VGG-16 (Simonyan & Zisserman, 2015) and ResNet20 (He et al., 2016) architectures of varying widths on the CIFAR-10 dataset. Results are presented in Figure 4.⁴

A clear relationship emerges between model width and linear mode connectivity, as measured by the loss barrier between solutions. Although $1 \times$ -sized models did not seem to exhibit linear mode connectivity, we found that larger width models decreased loss barriers all the way to zero. In Figure 4(right), we show what is to our knowledge the premiere demonstration of zero-barrier linear mode connectivity between two large ResNet models trained on a non-trivial dataset.

We highlight that relatively thin models do not seem to obey linear mode connectivity yet still exhibit similarities in training dynamics. This suggests that either our permutation selection methods are failing to find satisfactory permutations on thinner models or that some form of invariance other than permutation symmetries must be at play in the thin model regime.

5.4 MODEL PATCHING, SPLIT DATA TRAINING, AND IMPROVED CALIBRATION

Inspired by work on finetuning (Wortsman et al., 2022a), model patching (Raffel, 2021), and federated learning (McMahan et al., 2017; Konečný et al., 2016a;b), we study whether it is possible to synergistically merge the weights of two models trained on disjoint datasets. Consider, for example, an organization with multiple (possibly biased) datasets separated for regulatory (e.g., GDPR) or privacy (e.g., on-device data) considerations. Models can be trained on each dataset individually, but training in aggregate is not feasible. Can we combine separately trained models so that the merged model performs well on the entirety of the data?

To address this question, we split the CIFAR-100 dataset (Krizhevsky, 2009) into two disjoint subsets: dataset A, containing 20% examples labelled 0-49 and 80% labelled 50-99, and dataset B, vice versa. ResNet20 models A and B were trained on their corresponding datasets.

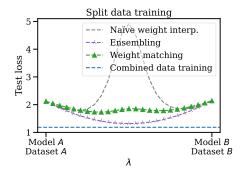


Figure 5: Models trained on disjoint datasets can be merged constructively. Algorithm 1 makes it possible for two ResNet models trained on disjoint, biased subsets of CIFAR-100 to be merged in weight space such that their combination outperforms both input models in terms of test loss on the combined dataset.

Figure 5 shows the result of merging the two models with weight matching. For comparison, we benchmark naïve weight interpolation, ensembling of the model logits, and full-data training.

As expected, merging separately trained models did not match the performance of an omniscient model trained on the full dataset or an ensemble of the two models with twice the number of effective weights. On the other hand, we did manage to merge the two models in weight space, achieving an interpolated model that outperforms both input models in terms of test loss while using half the memory and compute required for ensembling. Furthermore, the merged model's probability estimates are better calibrated than either of the input models as demonstrated in Figure 10. Algorithm 1 also vastly outperformed naïve interpolation, the status quo for model combination in federated learning and distributed training.

 $^{^4}$ Unfortunately, $8 \times$ width VGG-16 training was unattainable since it exhausted GPU memory on available hardware at the time of writing.

6 RELATED WORK

(Linear) mode connectivity. Freeman & Bruna (2017) introduced the concept of mode connectivity, i.e., that SGD solutions in the loss landscape are connected by constant-loss curves in weight space. Further explorations were undertaken in Garipov et al. (2018); Draxler et al. (2018), among others. Tatro et al. (2020) explored non-linear mode connectivity modulo permutation symmetries. Frankle et al. (2020) demonstrated a connection between *linear* mode connectivity and the lottery ticket hypothesis. Juneja et al. (2022) demonstrated that LMC does not always hold, even when fine-tuning. Hecht-Nielsen (1990); Chen et al. (1993) noted the existence of permutation symmetries, and Brea et al. (2019) implicated them as a source of saddle points in the loss landscape. Recently, the visionary work of Entezari et al. (2021) conjectured that SGD solutions could be linear mode connected modulo permutation symmetries and offered a battery of experiments buttressing this claim. Unlike previous works on LMC we accomplish zero-barrier paths between two independently-trained models with an algorithm that runs on the order of seconds.

Loss landscapes and training dynamics. Li et al. (2016); Yosinski et al. (2014) investigated whether independently trained networks learn similar features, and to what extent they transfer. Jiang et al. (2021) argued that independently trained networks meaningfully differ in the features they learn in certain scenarios. Zhang et al. (2019) studied the relative importance of layers. Benton et al. (2021) argued that SGD solutions form a connected volume of low loss. Pittorino et al. (2022) proposed a toroidal topology of solutions and a set of algorithms for symmetry removal. On the theoretical front, Kawaguchi (2016) proved that deep linear networks contain no local minima. Boursier et al. (2022); Chizat & Bach (2018); Mei et al. (2018) characterized the training dynamics of one-hidden layer networks, proving that they converge to zero loss. Godfrey et al. (2022); Simsek et al. (2021) investigated the algebraic structure of symmetries in neural networks and how this structure manifests in loss landscape geometry.

Federated learning and model merging. McMahan et al. (2017); Konečný et al. (2016a;b) introduced the concept of "federated learning," i.e., learning split across across multiple devices and datasets. Wang et al. (2020) proposed an exciting federated learning method in which model averaging is done after permuting units. Unlike this work, however, they proposed merging smaller "child" models into a larger "main" model, and doing so with a more specific, layer-wise matching algorithm that does not support residual connections or normalization layers. Raffel (2021); Matena & Raffel (2021); Sung et al. (2021) conceptualized the study of "model patching," i.e., the idea that models should be easy to modify and submit changes to. Ilharco et al. (2022) investigated model patching for the fine-tuning of open-vocabulary vision models. Singh & Jaggi (2020) proposed merging models by soft-aligning associations weights, inspired by optimal transport. Tatro et al. (2020); Liu et al. (2022) further explored merging models taking possible permutations into account. Wortsman et al. (2022a) demonstrated state-of-the-art ImageNet performance by averaging weights of models fine-tuned starting from some initial trained state.

7 DISCUSSION AND FUTURE WORK

We explore the role of permutation symmetries in the linear mode connectivity of SGD solutions. We introduce three novel algorithms to canonicalize independent neural network weights in order to make the loss landscape between them as flat as possible. In contrast to prior work, we linearly mode connect large ResNet models with no barrier in seconds to minutes. Despite presenting successes across multiple architectures and datasets, linear mode connectivity between thin models remains elusive. Therefore, we conjecture that permutation symmetries are a necessary piece, though not a complete picture, of the fundamental invariances at play in neural network training dynamics. In particular, we hypothesize that linear, possibly non-permutation, relationships connect the layerwise activations between models trained by SGD. In the infinite width limit, there exist satisfactory linear relationships that are also permutations.

An expanded theory and empirical exploration of other invariances – such as cross-layer scaling or general linear relationships between activations – presents an intriguing avenue for future work. Ultimately, we anticipate that a lucid understanding of loss landscape geometry will not only advance the theory of deep learning but will also promote the development of better optimization, federated learning, and ensembling techniques.

ETHICS STATEMENT

Merging models raises interesting ethical and technical questions about the resulting models. Do they inherit the same biases as their input models? Are rare examples forgotten when merging? Is it possible to gerrymander a subset of the dataset by splitting its elements across many shards?

Deployment of any form of model merging ought to be paired with thorough auditing of the resulting model, investigating in particular whether the merged model is representative of the entirety of the data distribution.

REPRODUCIBILITY STATEMENT

Our code is open sourced at https://github.com/REDACTED/REDACTED. Our experimental logs and downloadable model checkpoints are fully open source at https://wandb.ai/REDACTED/REDACTED.

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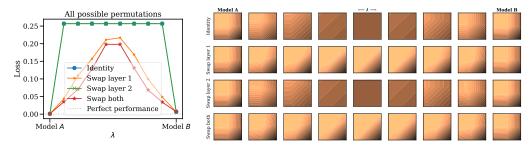


Figure 6: A counterexample. There exist models such that no possible permutation of weights allows for linear mode connectivity. *Left:* performance of all possible linear interpolations between the two models. *Right:* A visualization of the prediction functions f(x) through each linear sweep. Each row corresponds to one of the four possible permutations, and each column corresponds to a value of λ , the linear interpolant. The existence of such cases suggests that linear mode connectivity is an artifact of SGD.

A APPENDIX

A.1 KNOWN FAILURE MODES

We emphasize that none of the techniques presented in this paper are silver bullets. Here we list the failure cases that the authors are presently aware of,

- 1. Models of insufficient width
- 2. Models in the initial stages of training
- 3. VGGs on MNIST
- 4. MNIST MLPs trained with SGD and too low of a learning rate (As found by REDACTED)
- 5. MNIST MLPs trained with Adam and too high of a learning rate (As found by REDACTED)

Furthermore, we believe other failure modes certainly exist but have yet to be discovered.

We are excited by the prospect of future work investigating these failure modes and improving our understanding of when and why model merging modulo permutation symmetries is feasible.

A.2 ADDITIONAL INFORMATION ON ALGORITHM 1

On currently available hardware (p3.2xlarge AWS instance with a NVIDIA V100 GPU), we observed the following timing results with Algorithm 1,

- 1. MLP (3 layers, 512 units each): 3 seconds
- 2. ResNet50 (1× width): 33 seconds
- 3. ResNet20 (32× width): 194 seconds

In addition, we tested the ability of Algorithm 1 to recover a known, randomly selected permutation. In a handful of experiments we found that Algorithm 1 was able to exactly recover the known, random permutation in just 3-4 of passes over the layers.

A.3 COUNTEREXAMPLE DETAILS

Consider a simple 2-dimensional classification task. Our data points are drawn $x \sim \text{Uniform}([-1,1]^2)$ and $y = \mathbf{1}_{x_1 < 0 \text{ and } x_2 > 0}$. Figure 7 provides a visualization of a sample of such data.

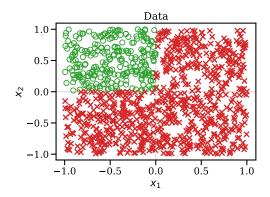


Figure 7: The counterexample classification problem data.

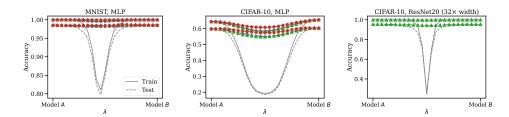


Figure 8: Top-1 accuracy results for the MNIST and CIFAR-10 models of Figure 2.

We utilize an MLP architecture consisting of two hidden layers, with two units each, and ReLU nonlinearities. Consider two weight assignments that both achieve a perfect fit to the data:

$$f_A(\boldsymbol{x}) = \begin{bmatrix} -1 & -1 \end{bmatrix} \sigma \begin{pmatrix} \begin{bmatrix} -1 & 0 \\ 0 & 1 \end{bmatrix} \sigma \begin{pmatrix} \begin{bmatrix} -1 & 0 \\ 0 & -1 \end{bmatrix} \boldsymbol{x} + \begin{bmatrix} 1 \\ 0 \end{bmatrix} \end{pmatrix} + \begin{bmatrix} 1 \\ 0 \end{bmatrix}$$
 (4)

$$f_B(\boldsymbol{x}) = \begin{bmatrix} -1 & -1 \end{bmatrix} \sigma \begin{pmatrix} \begin{bmatrix} 1 & 0 \\ 0 & -1 \end{bmatrix} \sigma \begin{pmatrix} \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} \boldsymbol{x} + \begin{bmatrix} 0 \\ 1 \end{bmatrix} \end{pmatrix} + \begin{bmatrix} 0 \\ 1 \end{bmatrix} \right). \tag{5}$$

We predict a positive label when $f(x) \ge 0$ and a negative label otherwise. Intuitively, these networks are organized such that each layer makes a classification whether $x_1 < 0$ or $x_2 > 0$. In model A, the first layer tests whether $x_2 > 0$, and the second layer tests whether $x_1 < 0$, whereas in model B the order is reversed. With a bit of algebra, it is possible to see that both f_A and f_B achieve perfect performance.

A.4 AUXILIARY PLOTS

A.5 STRAIGHT-THROUGH ESTIMATOR DETAILS

See Algorithm 2 for a complete description of the straight-through estimator algorithm.

A.6 MERGING MANY MODELS

We propose Algorithm 3 to merge the weights of more than two models at a time.

Following an argument similar to Lemma 2, it can be seen that Algorithm 3 terminates.

In our limited testing, we found that this algorithm converges quickly to solutions that extrapolate better than individual models and results in a merged model with better probability estimate calibration than any of the input models. For example, we present the results of this algorithm on MLPs trained on MNIST in Table A.6.

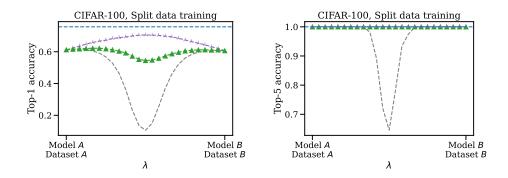


Figure 9: Accuracy results for the CIFAR-100 split data experiment.

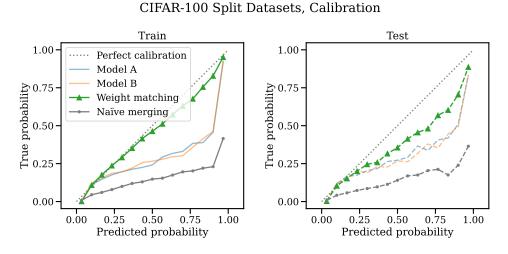


Figure 10: **Merging CIFAR-100 split data models results in superior probability calibration.** Although our merged model is not competitive in terms of top-1 accuracy in the CIFAR-100 split data experiment, we find that it has far better calibrated probability estimates than either of the input models.

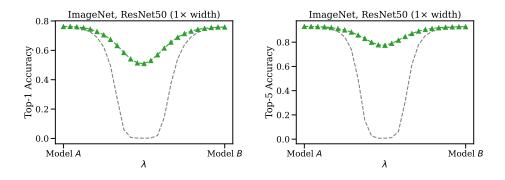


Figure 11: Accuracy results for merged ResNet50 (1× width) models on ImageNet.

Algorithm 2: Straight-through estimator training

Given: Model weights Θ_A , Θ_B , and a learning rate η .

Result: A permutation π of Θ_B such that $\mathcal{L}(\frac{1}{2}(\Theta_A + \pi(\Theta_B)))$ is approximately minimized.

Algorithm 3: MERGEMANY

```
Given: Model weights \Theta_1, \dots, \Theta_N
Result: A merged set of parameters \tilde{\Theta}.
```

In addition, we found that merging multiple models helps to calibrate the resulting model predictions. We present this effect in Figure 12.

A.7 FAILED IDEA: A METHOD FOR STEEPEST DESCENT

Imagine standing in weight space at Θ_A and trying to decide in which immediate direction to move in order to approach a Θ_B -equivalent point. There are many, many possible permutations of Θ_B – call them $\pi^{(1)}(\Theta_B), \pi^{(2)}(\Theta_B), \ldots$ – to aim for in the distance. Assuming that the loss landscape is in fact (quasi-)convex modulo these permutation symmetries, a natural choice would be to pick the $\pi^{(i)}(\Theta_B)$ that corresponds to the direction of steepest descent starting from Θ_A since we expect

| | Train Loss | Train Acc. | Test Loss | Test Acc. |
|-----------|------------|------------|-----------|-----------|
| Seed 1 | 0.0000 | 1.0000 | 0.1153 | 0.9856 |
| Seed 2 | 0.0000 | 1.0000 | 0.1531 | 0.9854 |
| Seed 3 | 0.0000 | 1.0000 | 0.1229 | 0.9855 |
| Seed 4 | 0.0000 | 1.0000 | 0.1108 | 0.9865 |
| Seed 5 | 0.0000 | 1.0000 | 0.1443 | 0.9871 |
| MERGEMANY | 0.0141 | 0.9952 | 0.0727 | 0.9831 |

Table 2: Merging multiple models decreases test loss by 43%. We train five separate MLPs on MNIST. Using Algorithm 3 we merge all these models together simultaneously. This produces a model that appears to be better calibrated than any of the input models, with superior test loss performance. We are excited by potential applications of this methodology in federated learning and ensembling, esp. along the lines of "model soups" (Wortsman et al., 2022a).

MergeMany Calibration

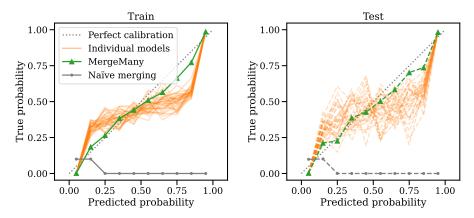


Figure 12: **Merging multiple models results in superior calibration.** Here we show the results of running Algorithm 3 on 32 MLP models trained on MNIST, with each model given access to a random 50% of the training dataset. The resulting merged model demonstrates substantively improved calibration of probability estimates on both the training and test datasets.

 $\pi^{(i)}(\Theta_B)$ to lie in the same basin as Θ_A . In other words,

$$\min_{\pi} \left. \frac{d\mathcal{L}(\Theta_A + \lambda(\pi(\Theta_B) - \Theta_A))}{d\lambda} \right|_{\lambda=0} = \min_{\pi} \left. \nabla \mathcal{L}(\Theta_A)^{\top} (\pi(\Theta_B) - \Theta_A) \right. \tag{6}$$

$$= -\nabla \mathcal{L}(\Theta_A)^{\top} \Theta_A + \min_{\pi} \left. \nabla \mathcal{L}(\Theta_A)^{\top} \pi(\Theta_B) \right. \tag{7}$$

Now, we are tenuously in a favorable situation: $\nabla \mathcal{L}(\Theta_A)$ is straightforward to compute, and picking the best π reduces to a matching problem. In particular it is a SOBLAP matching problem of the same form as in Section 3.2. In addition, there is a fast, exact solution for the single intermediate layer case (L=2).

In practice, we found that this method can certainly find directions of steepest descent, but that they are accompanied by high barriers in between the initial dip and $\pi(\Theta_B)$.

A.8 PROOF OF LEMMA 1

To lighten notation we use $\langle \cdot, \cdot \rangle = \langle \cdot, \cdot \rangle_F$ in this section.

Lemma. Given $A, B \in \mathbb{R}^{d \times d}$,

$$\min_{oldsymbol{P},oldsymbol{Q} ext{ perm. matrices}} \left\langle oldsymbol{P}oldsymbol{A}oldsymbol{Q}^ op,oldsymbol{B}
ight
angle$$

is strongly NP-hard and has no PTAS.

Proof. We proceed by reduction from the quadratic assignment problem (QAP) (Koopmans & Beckmann, 1957; Cela, 2013). Consider a QAP,

$$\min_{oldsymbol{P} ext{ perm. matrix}} \left\langle oldsymbol{P} oldsymbol{C} oldsymbol{P}^ op, oldsymbol{D}
ight
angle$$

for C, $D \in \mathbb{R}^{d \times d}$.

Now, pick $A = C + \lambda I$, $B = D - \lambda I$. The we have,

$$\min_{\boldsymbol{P},\boldsymbol{Q}} \langle \boldsymbol{P}(\boldsymbol{C} + \lambda \boldsymbol{I}) \boldsymbol{Q}^{\top}, \boldsymbol{D} - \lambda \boldsymbol{I} \rangle = \langle \boldsymbol{P} \boldsymbol{C} \boldsymbol{Q}^{\top} + \lambda \boldsymbol{P} \boldsymbol{Q}^{\top}, \boldsymbol{D} - \lambda \boldsymbol{I} \rangle \tag{8}$$

$$= \langle \boldsymbol{P} \boldsymbol{C} \boldsymbol{Q}^{\top}, \boldsymbol{D} \rangle - \lambda \langle \boldsymbol{P} \boldsymbol{C} \boldsymbol{Q}^{\top}, \boldsymbol{I} \rangle + \lambda \langle \boldsymbol{P} \boldsymbol{Q}^{\top}, \boldsymbol{D} \rangle - \lambda^{2} \langle \boldsymbol{P} \boldsymbol{Q}^{\top}, \boldsymbol{I} \rangle$$

$$= \langle \boldsymbol{P} \boldsymbol{C} \boldsymbol{Q}^{\top}, \boldsymbol{D} \rangle - \lambda \langle \boldsymbol{P}^{\top} \boldsymbol{Q}, \boldsymbol{C} \rangle + \lambda \langle \boldsymbol{P} \boldsymbol{Q}^{\top}, \boldsymbol{D} \rangle - \lambda^{2} \operatorname{tr}(\boldsymbol{P} \boldsymbol{Q}^{\top})$$
(10)

For sufficiently large λ , the $\operatorname{tr}(\boldsymbol{P}\boldsymbol{Q}^{\top})$ term will dominate. Letting $\alpha = \max\left(\max_{i,j}|C_{i,j}|,\max_{i,j}|D_{i,j}|\right)$, we can bound the other terms,

$$-d^{2}\alpha^{2} \leq \langle PCQ^{\top}, D \rangle \leq d^{2}\alpha^{2}$$
(11)

$$-\lambda d\alpha \le -\lambda \langle \boldsymbol{P}^{\top} \boldsymbol{Q}, \boldsymbol{C} \rangle \le \lambda d\alpha \tag{12}$$

$$-\lambda d\alpha \le \lambda \langle \boldsymbol{P} \boldsymbol{Q}^{\top}, \boldsymbol{D} \rangle \le \lambda d\alpha \tag{13}$$

Now there are two classes of solutions: those where P=Q and those where $P\neq Q$. We seek to make the best (lowest) possible $P\neq Q$ solution to have worse (higher) objective value than the worst (highest) P=Q solution. When P=Q, the highest possible objective value is

$$d^2\alpha^2 + \lambda d\alpha + \lambda d\alpha - \lambda^2 d$$

and similarly, the lowest possible objective value when $oldsymbol{P}
eq oldsymbol{Q}$ is

$$-d^2\alpha^2 - \lambda d\alpha - \lambda d\alpha - \lambda^2 d + \lambda^2$$

where the final term is due to the fact that at least one entry of PQ^{\top} must be 0. With some algebra, it can be seen that $\lambda > 5d\alpha$ is sufficient to guarantee that all P = Q solutions are superior to all $P \neq Q$ solutions.

Now when P = Q, all frivolous terms reduce to constants and we are left with the QAP objective:

$$\begin{split} \min_{\boldsymbol{P}} \ \langle \boldsymbol{P}\boldsymbol{C}\boldsymbol{P}^{\top}, \boldsymbol{D} \rangle - \lambda \langle \boldsymbol{P}^{\top}\boldsymbol{P}, \boldsymbol{C} \rangle + \lambda \langle \boldsymbol{P}\boldsymbol{P}^{\top}, \boldsymbol{D} \rangle - \lambda^{2} \operatorname{tr}(\boldsymbol{P}\boldsymbol{P}^{\top}) \\ = -\lambda \operatorname{tr}(\boldsymbol{C}) + \lambda \operatorname{tr}(\boldsymbol{D}) - \lambda^{2} d + \min_{\boldsymbol{P}} \ \langle \boldsymbol{P}\boldsymbol{C}\boldsymbol{P}^{\top}, \boldsymbol{D} \rangle \end{split}$$

completing the reduction. QAP is known to be strongly NP-hard (Koopmans & Beckmann, 1957; Sahni & Gonzalez, 1976) and MaxQAP is known to not admit any PTAS (Makarychev et al., 2014), thus completing the proof.

A.9 PROOF OF LEMMA 2

Lemma. Algorithm 1 terminates.

Proof. We proceed by contradiction.

Consider a graph with each possible permutation $\pi_i = \{P_1, \dots, P_{L-1}\}$ as a vertex and directed edges $\pi_i \to \pi_j$ if π_j can be reached from π_i with a single P_ℓ update, as in Algorithm 1. (Ignore those updates that result in no change to P_ℓ in order to avoid $\pi_i \to \pi_i$ cycles.) Let $\rho(\pi) = \text{vec}(\Theta_A) \cdot \text{vec}(\pi(\Theta_B))$ denote the utility of a particular π . Note that $\pi_i \to \pi_j$ implies $\rho(\pi_i) < \rho(\pi_j)$. There exist finitely many possible permutations π_i , meaning that a failure to terminate must involve a cycle in the graph $\pi_1 \to \cdots \to \pi_n \to \pi_1$. However ρ forms a total order on the vertices and therefore we have a contradiction.

A.10 EXTENDED RELATED WORK

Differentiating through permutations. Akin to differentiable permutation learning, many prior works have studied differentiable sorting (Grover et al., 2019; Prillo & Eisenschlos, 2020; Cuturi et al., 2019; Petersen et al., 2022; 2021; Mena et al., 2018). Blondel et al. (2020) studied differentiable sorting and ranking with asymptotics that correspond to their non-differentiable versions. Fogel et al. (2015) explored recovering the linear orderings of items based on pairwise information, another form of permutation optimization. Bengio et al. (2013) introduced the straight-through estimator for differentiating through discrete projections that we utilize in Section 3.3.