

Unexplainable Explanations: Towards Interpreting tSNE and UMAP Embeddings

Andrew Draganov^{1*} and Simon Dohn¹

^{1*}Department of Computer Science, Aarhus University, Denmark.

*Corresponding author(s). E-mail(s): draganovandrew@cs.au.dk;
Contributing authors: dohn@cs.au.dk;

Abstract

It has become standard to explain neural network latent spaces with attraction/repulsion dimensionality reduction (ARDR) methods like tSNE and UMAP. This relies on the premise that structure in the 2D representation is consistent with the structure in the model’s latent space. However, this is an unproven assumption – we are unaware of any convergence guarantees for ARDR algorithms. We work on closing this question by relating ARDR methods to classical dimensionality reduction techniques. Specifically, we show that one can fully recover a PCA embedding by applying attractions and repulsions onto a randomly initialized dataset. We also show that, with a small change, Locally Linear Embeddings (LLE) can reproduce ARDR embeddings. Finally, we formalize a series of conjectures that, if true, would allow one to attribute structure in the 2D embedding back to the input distribution.

Keywords: XAI, Dimensionality Reduction, tSNE, UMAP

1 Introduction

The modern machine learning engineer has no doubt used a neural network without fully knowing what the model chose to prioritize in the data. Indeed, deep learning models are known to take shortcuts, focusing on spurious correlations rather than semantic-level features in the data. Thus, the field of Explainable AI (XAI) has risen to prominence to help analyze machine learning models and align the user’s intuition with the model’s feature extraction process.

The first step towards interpreting what a network has learned is to inspect the distributions in its feature space. Since deep learning embeddings are too high-dimensional to interpret on their own, it is common to perform dimensionality reduction (DR) to get the embedding vectors into 2D. To this end, tSNE [1] and UMAP [2] have become the de-facto tool for visualizing learned representations due to their attractive properties – they are unsupervised, fast to optimize and produce 2D distributions with neatly separated clusters. In fact, it has become almost standard to show that a feature extractor works as intended by plotting its embeddings with tSNE or UMAP; a non-exhaustive list of examples can be found in [3–47]. While most of these examples are in the unsupervised learning setting, we point out that tSNE and UMAP are also used to augment understanding of supervised latent spaces.

However, despite tSNE and UMAP giving interpretable embeddings and being used for network explainability, they themselves are not well understood. For example, [48] showed that UMAP’s implementation does not optimize the intended loss function, implying that the theoretical motivation may not hold in practice. Furthermore, it has been a point of active research to investigate how the loss functions, gradient descent strategies, and heuristic accelerations alter the structure of tSNE and UMAP embeddings [49–53], with many works suggesting that the common wisdom surrounding tSNE and UMAP may not hold. Thus, it is unclear whether conclusions drawn from the 2D embeddings can be carried over to the neural network’s learned representation. Essentially, the community is trying to solve an explainability problem with a tool that has a well-studied explainability problem.

Luckily, modern methods like tSNE and UMAP exist in a landscape of otherwise well-understood DR techniques. For example, it is known that Principal Component Analysis (PCA) maximally preserves variance from the input distribution. Thus, given a PCA embedding, one can confidently attribute structure back to the original dataset [54]. Similar conclusions can be drawn for other classic DR techniques such as Locally Linear Embeddings (LLE) [55, 56], ISOMAP [57], Laplacian Eigenmaps [58], and Multi-Dimensional Scaling (MDS) [59], to name a few. Importantly, their explainability is a direct consequence of the convergence guarantees of each algorithm. Unfortunately, these do not provide the intuitive separation that deep learning practitioners want and are therefore not as popular for visualizing latent spaces. Thus, the field has been inadvertently partitioned: DR methods are either well-understood or are useful for deep learning explainability, but not both.

1.1 Our Contributions

Our work aims to bridge the gap between these two types of methods. We start by defining a framework for representing methods such as PCA and LLE in the tSNE/UMAP setting. Specifically, we show that one can provably obtain PCA embeddings by performing attractions and repulsions between points and that this is robust to fast low-rank approximations. Furthermore, minimizing the PCA and LLE objective functions with the UMAP kernel on the 2D embedding gives similar gradients as those found in tSNE/UMAP. We verify this experimentally by showing that minimizing the LLE objective with this kernel gives comparable embeddings to tSNE/UMAP. Thus, we show that classical methods are reproducible in the modern DR framework.



Fig. 1: UMAP/tSNE (top/bottom) embeddings of the last conv. layer of VGG11 [60] on MNIST mid-training. The columns represent different epochs of training. 3 columns have $>90\%$ accuracy while one has 72% accuracy. We purposefully omit which column corresponds to the low-accuracy latent space.

This naturally leads us to ask whether the relationship goes the other way. Specifically, we conjecture that the tSNE/UMAP gradient descent heuristics minimize the LLE objective with two kernels. If this holds true, it would suggest that one does not require gradient descent to find an embedding that is provably similar to those obtained by tSNE/UMAP. In this sense, we hope to facilitate future work towards the open question: “Which properties in the tSNE/UMAP embedding guarantee structure in the original dataset?” We identify that the crux of this question lies in the poorly-studied dynamics imposed by the kernel added on the embedding and discuss opportunities for future work therein.

2 Related Work

There has been a significant attempt in recent years to explain deep learning models’ learned representations. Some methods attempt to explain a model’s decision-making using specific data examples, for instance through saliency maps [61–63] or counterfactual explanations [64, 65]. Other approaches focus on visualizing some aspect of the network itself, such as feature activations, convolution filters [63], and the feature vectors by finding neighbors [66], or embedding them into 2D, as we will discuss below.

For the rest of the paper we will refer to the class of gradient-based dimensionality reduction techniques that includes tSNE and UMAP as Attraction/Repulsion DR (ARDR) methods. The common theme among these is that they define notions of similarity in the input \mathbf{X} and the embedding \mathbf{Y} and minimize a loss function by attracting points in \mathbf{Y} that should be similar and repelling points that should be dissimilar. A non-exhaustive list of methods includes tSNE [1, 67], UMAP [2], ForceAtlas2 [68], LargeVis [69] and PacMAP [50]. Similar methods such as TriMAP [70] discuss this in the context of triplets but we note that the underlying schema of gradient descent by attractions and repulsions remains the same.

Despite the popularity of ARDR methods, there has been a recent wave of literature that disputes the common wisdom surrounding them. We recommend [71] to increase intuition regarding how tSNE’s hyperparameters affect one’s ability to relate the structures in the input and 2D embedding. We note that UMAP has similar issues to those raised for tSNE in [71]. For more rigorous analysis that dispels myths regarding ARDR methods, [49] showed that UMAP’s seemingly stronger attractions are a result of the sampling strategy rather than the topological properties of the algorithms. This was followed by [48], which noted the surprising fact that UMAP does not actually optimize its stated loss function due to the heuristics it employs during gradient descent. Further analysis of ARDR loss functions can be found in [50, 51]. We note that [52] found that one can replace the KL divergence in UMAP by the sum of squared errors without impacting the embeddings in practice. Furthermore, the manifold learning intuitions regarding ARDR methods were questioned in [53, 70, 72], where it was shown that it is unclear whether tSNE and UMAP are preserving local/global structure in the expected manner. Lastly, other works have sought to unify tSNE and UMAP by verifying that they are effectively identical up to a hyperparameter [52] and can both be reproduced within the contrastive learning framework [73].

We are less aware of literature comparing classical DR techniques to the modern wave of gradient-based ones. [51] defines a framework to unify classical and modern DR methods, but the generality of the approach makes it difficult to make direct claims regarding the connection between methods like PCA and UMAP. Furthermore, tSNE does not fit into their framework in a natural manner. Indeed, the reference most similar to ours is [53], where the authors analyze the manifold-learning properties of popular DR approaches towards the goal of improved explainability. They study how well various DR techniques preserve locality via a novel measure and show that (1) tSNE/UMAP preserve locality better than most classical methods¹, and (2) that tSNE/UMAP distort the Euclidean relationships more strongly than other DR methods. We share the authors’ surprise regarding the lack of literature on explainable DR methods since dimensionality reduction is a natural step towards gathering intuition on learned representations.

We approach the question of explainability from the other direction. Whereas [53] studies the experimental properties of embeddings obtained by modern and classical methods, we instead relate their theoretical foundations. Specifically, Section 3 provides background on PCA, LLE, and ARDR methods, after which Sections 4 and 5 show how to interpret classical methods in terms of attractions and repulsions between points. We conclude by motivating conjectures and open questions in 6.

3 Preliminaries

3.1 Principal Component Analysis (PCA)

Likely the most famous dimensionality reduction algorithm, PCA finds the orthogonal basis in $\mathbb{R}^{n \times d}$ that maximally preserves the variance in the centered dataset [74]. Let $\mathbf{CX} = \mathbf{U}_X \mathbf{\Sigma}_X \mathbf{V}_X^\top$ be the singular value decomposition (SVD) of the centered dataset,

¹Although there are situations where PCA performs best.

where $\mathbf{C} = \mathbf{I} - \frac{1}{n}\mathbf{J}$ is the centering matrix with \mathbf{J} the all-1 matrix. This gives us an expression for the positive semidefinite (psd) Gram matrix $\mathbf{C}\mathbf{G}_X\mathbf{C} = \mathbf{C}\mathbf{X}\mathbf{X}^\top\mathbf{C} = \mathbf{U}_X\mathbf{\Sigma}_X^2\mathbf{U}_X^\top$. The principal components of $\mathbf{C}\mathbf{X}$ are defined as $\mathbf{U}_X\mathbf{\Sigma}_X$ can therefore be found via eigen-decomposition on $\mathbf{C}\mathbf{G}_X\mathbf{C}$.

This has a natural extension to kernel methods [75]. Suppose that the kernel function $k_x(x_i, x_j) = \langle \phi(x_i), \phi(x_j) \rangle$ corresponds to an inner product space. Then the kernel matrix $[\mathbf{K}]_{ij} = k(x_i, x_j)$ is a psd Gram matrix for the space defined by ϕ and admits an equivalent procedure for finding principal components as traditional PCA.

3.2 Locally Linear Embedding (LLE)

Since PCA embeddings are obtained via linear projection operations, one can think of them as preserving large distances in the dataset. On the other hand, LLE preserves local neighborhoods by finding the \mathbf{Y} that maximally recreates linear combinations of nearest-neighbors in \mathbf{X} . Assume that we have the k -nearest-neighbor graph on \mathbf{X} and let $K_i = \{e_{i1}, \dots, e_{ik}\}$ represent the indices of x_i 's k nearest neighbors in \mathbf{X} . LLE then proceeds by finding the $\mathbf{W} \in \mathbb{R}^{n \times n}$ such that

$$\sum_i \|x_i - \sum_{j \in K_i} w_{ij}x_j\|_2^2 = \|\mathbf{X} - \mathbf{W}\mathbf{X}\|_F^2 = \|(\mathbf{I} - \mathbf{W})\mathbf{X}\|_F^2$$

is minimized under the constraint $\sum_{j \in K_i} w_{ij} = 1$ for all i . That is, we find the weights such that each point x_i is represented as a linear combination of its neighbors. This implies that the i -th row of \mathbf{W} must be zero on the indices l where x_l is not x_i 's nearest neighbor. It can be shown that such a \mathbf{W} always exists and each row can be found by solving an eigenproblem [55, 56].

Having found the \mathbf{W} that represents local neighborhood relationships in \mathbf{X} , step 2 of LLE then finds the $\mathbf{Y} \in \mathbb{R}^{n \times d}$ such that

$$\sum_i \|y_i - \sum_{j \in K_i} w_{ij}y_j\|_2^2 = \|\mathbf{Y} - \mathbf{W}\mathbf{Y}\|_F^2$$

is minimized², subject to the constraint that the columns of \mathbf{Y} form an orthogonal basis, i.e. $\frac{1}{n}\mathbf{Y}^\top\mathbf{Y} = \mathbf{I}$. The embedding \mathbf{Y} is given by the eigenvectors of $\mathbf{M} = (\mathbf{I} - \mathbf{W})^\top(\mathbf{I} - \mathbf{W})$ that correspond to the smallest d positive eigenvalues (Derivation in B.1).

LLE can again be combined with kernels on \mathbf{X} and \mathbf{Y} to perform more sophisticated similarity calculations [56]. Let \mathbf{K}_X be defined as in Section 3.1 and let $k_y(y_i, y_j) = \langle \psi(y_i), \psi(y_j) \rangle$ be a psd kernel that defines a corresponding $\mathbf{K}_Y \in \mathbb{R}^{n \times n}$ with $[\mathbf{K}_Y]_{ij} = k_y(y_i, y_j)$. Then we can define the objective function during the first step as finding the \mathbf{W} that minimizes $\text{Tr}((\mathbf{I} - \mathbf{W})\mathbf{K}_X(\mathbf{I} - \mathbf{W})^\top)$ such that $\sum_j w_{ij} = 1$ for all i . Similarly, the second step under a kernel function \mathbf{K}_Y seeks the \mathbf{Y} such that $\text{Tr}((\mathbf{I} - \mathbf{W})\mathbf{K}_Y(\mathbf{I} - \mathbf{W})^\top)$ is minimized.

²Note that the \mathbf{W} and K_i are the same as in the first step.

3.3 Gradient Dimensionality Reduction Methods

We now switch gears from the directly solvable methods to the gradient-based ones which have risen to prominence in the last decade. This includes algorithms such as UMAP, tSNE, LargeVis, and PacMAP, just to name a few.

3.3.1 In theory

ARRDR methods

Assume that we are given an input $\mathbf{X} \in \mathbb{R}^{n \times D}$, a randomly initialized embedding $\mathbf{Y} \in \mathbb{R}^{n \times d}$ and two psd kernel functions $k_x(x_i, x_j) = \langle \phi(x_i), \phi(x_j) \rangle$ and $k_y(y_i, y_j) = \langle \psi(y_i), \psi(y_j) \rangle$. These then define psd matrices $\mathbf{K}_X, \mathbf{K}_Y \in \mathbb{R}^{n \times n}$, where each (i, j) -th entry represents the set of per-point similarities in \mathbf{X} and \mathbf{Y} respectively. Given this setup, the goal of gradient-based DR methods is to find the embedding \mathbf{Y} such that \mathbf{K}_Y is maximally similar to \mathbf{K}_X with respect to some matrix-wise loss function.

For the remainder of this section, the reader can assume that k_x is the exponential RBF kernel and k_y is the quadratic Cauchy kernel; these are the common choices across ARDR methods. However, we note that the upcoming generalizations only require that k_y be a function of the squared Euclidean distance. We will write $k_y(\|y_i - y_j\|_2^2)$ when emphasizing this point.

Since the kernels are usually chosen such that k_x and k_y are necessarily in $[0, 1]$, the matrices can be treated as probability distributions³. Thus, the question of how well \mathbf{K}_Y represents \mathbf{K}_X is traditionally quantified using the KL-divergence $KL(\mathbf{K}_X || \mathbf{K}_Y)$. For example, tSNE and UMAP state the following loss functions⁴

$$\mathcal{L}_{tSNE}(\mathbf{X}, \mathbf{Y}) = \sum_{\substack{i,j=1 \\ i \neq j}}^n k_x(x_i, x_j) \log \left(\frac{k_x(x_i, x_j)}{k_y(y_i, y_j)} \right) \quad (1)$$

$$\mathcal{L}_{UMAP}(\mathbf{X}, \mathbf{Y}) = \sum_{\substack{i,j=1 \\ i \neq j}}^n k_x(x_i, x_j) \log \left(\frac{k_x(x_i, x_j)}{k_y(y_i, y_j)} \right) + (1 - k_x(x_i, x_j)) \log \left(\frac{1 - k_x(x_i, x_j)}{1 - k_y(y_i, y_j)} \right). \quad (2)$$

By then evaluating the gradient with respect to y_i , one finds that it can be written as a set of attractive and repulsive terms

$$\nabla_{y_i} = -c \sum_j (\mathcal{A}_{ij} - \mathcal{R}_{ij}) \cdot \frac{d k_y(\|y_i - y_j\|_2^2)}{d \|y_i - y_j\|_2^2} \cdot (y_i - y_j)$$

where \mathcal{A}_{ij} and \mathcal{R}_{ij} act as scalars on the vector $(y_i - y_j)$ and the $\frac{d k_y(\|y_i - y_j\|_2^2)}{d \|y_i - y_j\|_2^2}$ scalar comes as a result of the chain rule over k_y . In this sense, we have that y_i is attracted to y_j according to \mathcal{A}_{ij} and repelled according to \mathcal{R}_{ij} . Performing gradient descent then

³Either as a matrix of n^2 Bernoulli random variables as in UMAP or, if the matrix sums to 1, as a single probability distribution as in tSNE

⁴We ignore the iterands where $i = j$ because it is assumed that a point should have no effect on itself within the embedding.

means that we apply these attractions and repulsions across all n^2 pairs of points, where each force’s strength is a function of k_x , k_y , and the length of the vector $y_i - y_j$.

Generalization

This attraction/repulsion framework is not restricted to the gradient of the KL-divergence, however. Indeed, the $(y_i - y_j)$ vectors come directly from the fact that k_y is a function of the squared Euclidean distance. This is evidenced by the chain rule, as any derivative of $k_y(\|y_i - y_j\|_2^2)$ with respect to y_i must depend on the derivative of $\|y_i - y_j\|_2^2$. To this end, we see

$$\begin{aligned} \|y_i - y_j\|_2^2 &= y_i^\top y_i - 2y_i^\top y_j + y_j^\top y_j \\ \implies \frac{\partial \|y_i - y_j\|_2^2}{\partial y_i} &= 2(y_i - y_j), \end{aligned}$$

implying that the gradient of $k_y(\|y_i - y_j\|_2^2)$ will necessarily be some scalar acting on $(y_i - y_j)$. Furthermore, any loss function of the following form will incur gradients with attractions and/or repulsions between points.

$$\mathcal{L}_{\mathcal{F}}(\mathbf{X}, \mathbf{Y}) = \sum_{\substack{i,j=1 \\ i \neq j}}^n \mathcal{F}(k_x(x_i, x_j), k_y(y_i, y_j)) \tag{3}$$

where \mathcal{F} is any function that grows as $k_x(x_i, x_j)$ and $k_y(y_i, y_j)$ diverge. The most natural such objective function is the Frobenius norm

$$\mathcal{L}_{Frob}(\mathbf{X}, \mathbf{Y}) = \|\mathbf{K}_X - \mathbf{K}_Y\|_F^2 = \sum_{\substack{i,j=1 \\ i \neq j}}^n (k_x(x_i, x_j) - k_y(y_i, y_j))^2.$$

and it was shown in [52] that substituting this for UMAP’s KL-divergence objective produces effectively equivalent embeddings (discussed further in Section 5.1).

We will use the term *attraction/repulsion framework* (A/R framework) to refer to an analysis that uses attractions and repulsions. We will refer to the task of finding an optimal such rank- d embedding as the *A/R objective*. That is, $\min_Y \mathcal{L}_{\mathcal{F}}(\mathbf{X}, \mathbf{Y})$.

3.3.2 In practice

Many gradient-based DR methods develop the above framework before delving into heuristics that approximate the solution. For example, a common trick is to notice that an exponential kernel for k_x is likely to be 0 for distant points. Since \mathcal{A}_{ij} linearly depends on $k_x(x_i, x_j)$ in most ARDR methods, the attractions are therefore weak if x_i and x_j are far away. To this end, the common heuristic is to compute a k -NN graph over \mathbf{X} and only calculate attractions along pairs of nearest neighbors. Furthermore, it has been shown that computing a subset of the repulsions acting on a point is equivalent in expectation to computing all of the repulsions.

Thus, the usual optimization strategy is to simulate gradient descent by iterating between the relevant attractions and sampled repulsions acting on each point. Specifically, one attracts y_i to the points corresponding to x_i 's k nearest neighbors in \mathbf{X} . Then one finds ck random points from which to repel y_i , where c is some appropriately chosen constant.

3.4 Next Steps

Our goal for this paper is to use the explainability inherent in PCA and LLE to present concrete interpretability for ARDR methods as a whole. To do this, we first show that methods like PCA and LLE are easily framed in terms of the A/R framework – their objective functions can be minimized by performing attractions and repulsions on points in a randomly initialized embedding. Furthermore, their gradients under a nonlinear k_y strongly resemble those in UMAP and other gradient-based methods.

4 PCA in the ARDR Framework

Representing a learning problem in the ARDR framework requires two things. First, we must find the k_x , k_y and \mathcal{F} such that the objective can be written in the form of Equation 3. Second, we must show that the minimum of the A/R formulation is indeed the optimum of the original problem. We show this for PCA as it is the most natural DR algorithm and therefore sets the stage for future steps.

4.1 PCA as Attractions and Repulsions

PCA is often presented as an SVD-based algorithm to find the optimal low-rank representation of \mathbf{CX} . By the Eckart-Young-Mirsky theorem, the optimal low-rank representation with respect to the Frobenius norm is given by the $\tilde{\mathbf{X}}$ s.t. $\|\mathbf{CX} - \tilde{\mathbf{X}}\|_F$ is minimized subject to $\text{rank}(\tilde{\mathbf{X}}) \leq d$. It is well-known that, for centered \mathbf{X} , this is given by the first d principal components of $\mathbf{CG}_X\mathbf{C} = \mathbf{CXX}^\top\mathbf{C}$, which is exactly the embedding \mathbf{Y} obtained by PCA. With this as justification, we state PCA's objective function as

$$\min_{\mathbf{Y}} \|\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\|_F^2 \quad \text{s.t.} \quad \mathbf{Y}^\top\mathbf{C}\mathbf{Y} = \mathbf{I}$$

where $\mathbf{G}_Y = \mathbf{Y}\mathbf{Y}^\top$. Note that the centering matrix \mathbf{C} is symmetric and idempotent, i.e. $\mathbf{C}^\top\mathbf{C} = \mathbf{C}^2 = \mathbf{C}$. Under this setting, we have that k_x and k_y are both the standard inner product and $\mathcal{F}(a, b) = (a - b)^2$. We now show that this is minimized if and only if \mathbf{Y} is the PCA projection of \mathbf{X} .

Lemma 1. The minimum of $L_{PCA}(\mathbf{X}, \mathbf{Y}) = \|\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\|_F^2$ is only obtained when \mathbf{Y} is the PCA projection of \mathbf{X} up to orthogonal transformation.

The proof is given in section A.1 of the supplementary material. We now provide the formula for the PCA gradient and verify that it can be represented as attractions and repulsions between points.

Corollary 1. The PCA gradient $\nabla_{PCA} \in \mathbb{R}^{n \times d}$ is given by

$$\nabla_{PCA} = -2\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C} : d(\mathbf{C}\mathbf{G}_Y\mathbf{C}) = -4\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\mathbf{Y}, \quad (4)$$

where the \cdot operator implies the Frobenius inner product.

We derive this in Section A.2 of the supplementary material. Furthermore, the next lemma states that this gradient can be written in terms of attractions and repulsions.

Lemma 2. Let $\mathbf{L} \in \mathbb{R}^{n \times n}$ be any matrix of the form $\mathbf{L} = \mathbf{C}\mathbf{A}\mathbf{C}$ for $\mathbf{A} \in \mathbb{R}^{n \times n}$ and let α be a constant. Then any gradient of the form $\nabla = \alpha\mathbf{L} : d(\mathbf{C}\mathbf{G}_Y\mathbf{C})$ can be expressed via attractions and repulsions on \mathbf{Y} . Furthermore, the A/R-style gradient acting on y_i has the form $\nabla_{y_i} = \alpha \sum_j l_{ij}(y_i - y_j)$.

The proof is given in Section A.3.

4.2 PCA Convergence

Given this gradient formulation, the remaining question is whether gradient descent will converge to the PCA embedding. Our next result shows this to be the case.

Theorem 1. Let \mathbf{Y} be any dataset in $\mathbb{R}^{n \times d}$ and let $\mathbf{L} = -4\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}$ for dataset $\mathbf{X} \in \mathbb{R}^{n \times D}$. Then there always exists a γ and $\mathbf{Y}' = (\mathbf{I} + \gamma\mathbf{L})\mathbf{Y}$ such that $\mathcal{L}_{PCA}(\mathbf{X}, \mathbf{Y}') \leq \mathcal{L}_{PCA}(\mathbf{X}, \mathbf{Y})$. Furthermore, continuously performing this gradient update will necessarily converge to the PCA embedding of \mathbf{X} .

The proof can be found in section A.4 of the supplementary material. Interestingly, this proof only requires that \mathbf{G}_X be a real, symmetric matrix. So not only does it trivially extend to Kernel PCA, it also holds for any real, symmetric similarity matrix, such as those described in [76]. We are thus guaranteed to obtain a PCA embedding of \mathbf{X} if we randomly initialize the pointset \mathbf{Y} and apply the gradient update in Equation 4. This naturally extends to PCA gradients in the A/R setting as well. Figure 2 shows the convergence of PCA by gradient descent on the MNIST dataset.

Performing this gradient descent is impractical, however, as one must calculate a new $\mathbb{R}^{n \times n} \times \mathbb{R}^{n \times d}$ matrix product at every epoch. In many cases, however, our similarity functions are psd and are therefore suitable for fast approximations. Given psd matrix \mathbf{A} and its SVD-based rank- k approximation \mathbf{A}^k , there are sublinear-time methods [77] for obtaining \mathbf{A}' such that

$$\|\mathbf{A} - \mathbf{A}'\|_F^2 \in (1 \pm \varepsilon)\|\mathbf{A} - \mathbf{A}^k\|_F^2 \quad (5)$$

Our next result shows that these approximations do not significantly affect the gradient descent convergence. Let ∇ be the full PCA gradient and ∇' be the gradient obtained using the approximation in Eq. 5.

Lemma 3. Let λ_{x_i} be the i -th eigenvalue of \mathbf{G}_X and let \mathbf{G}_X^k be the optimal rank- k approximation of \mathbf{G}_X . Then $\langle \nabla, \nabla' \rangle_F > 0$ as long as

$$\|\mathbf{G}_X - \mathbf{G}_Y\|_F^2 \geq (1 + \varepsilon) \frac{\lambda_{x_1}}{\lambda_{x_k}} \|\mathbf{G}_X - \mathbf{G}_X^k\|_F^2$$

We prove this in Section A.5 of the supplementary material. This effectively means that, as long as \mathbf{G}_Y is a worse⁵ approximation to \mathbf{G}_X than \mathbf{G}_X^k , the approximate gradient points in a similar direction to the true one. Notice that once this is not the case, then we have that $\|\mathbf{G}_X - \mathbf{G}_Y\|_F^2$ is an α -approximation of $\|\mathbf{G}_X - \mathbf{G}_X^k\|_F^2$, with

⁵Up to scaling by ε and the eigengap

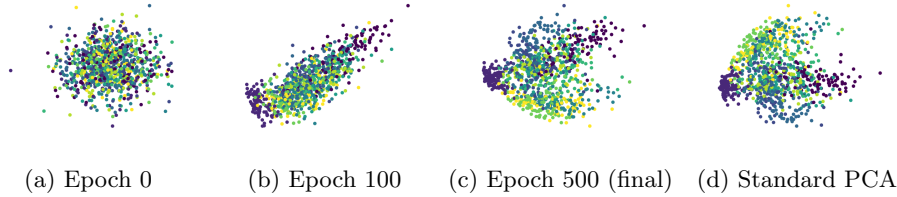


Fig. 2: PCA on 1000 MNIST samples by gradient descent. The final result is equivalent to the standard PCA embedding up to orthogonal transformation, i.e. reflection on x-axis.

$\alpha = (1 + \epsilon)(\lambda_{x_1}/\lambda_{x_k})$. We conjecture that this line of reasoning could lead to a fast method for approximating PCA that does not require an eigendecomposition – one simply initializes a \mathbf{Y} and performs the necessary gradient updates in sublinear time.

5 AR Problems with Two Kernels

Using the blueprint developed in sections 3 and 4, we now show that ARDR methods such as UMAP can be emulated using classical methods and gradient descent. The key observation is simply that ARDR methods have a kernel on \mathbf{Y} while classical methods do not. Unfortunately, this kernel is also what introduces much of the difficulty when analyzing convergence properties.

To see this, consider that the above theoretical results on PCA hold for the linear kernel⁶ on \mathbf{Y} but no longer hold as soon as k_y is non-linear. This is due to the fact that differentiating with respect to k_y induces a new scalar by the chain rule. Indeed, the AR problem $\mathcal{L}_{\mathcal{F}}(\mathbf{X}, \mathbf{Y})$ will necessarily have gradients acting on y_i of the form

$$\nabla_{y_i} = \sum_j l_{ij} \cdot \frac{d k_y(\|y_i - y_j\|_2^2)}{d \|y_i - y_j\|_2^2} \cdot (y_i - y_j), \quad (6)$$

where l_{ij} depends on k_x , k_y , and the chosen loss function \mathcal{F} . We derive Equation 6 in the proof of Lemma 4. Observe that if k_y is the centered linear kernel then the derivative term is constant. Note, if \mathcal{F} is the Frobenius norm, then $l_{ij} = [\mathbf{K}_X - \mathbf{K}_Y]_{ij}$.

We relied on these observations heavily when proving the results in Section 4. Specifically, for $\mathbf{K}_Y \sim \mathbf{G}_Y$, the optimal solution must have $\mathbf{K}_X - \mathbf{K}_Y$ of rank $D - d$ while \mathbf{Y} has rank d . Thus, $(\mathbf{K}_X - \mathbf{K}_Y)\mathbf{Y}$ must be zero by orthogonality. However, a non-linear k_y impedes this line of reasoning. First, it is unclear what dynamics the non-constant derivative term in Eq. 6 introduces. Second, the above orthogonality argument is broken for non-linear k_y – we could have that $\text{rank}(\mathbf{K}_Y) > d$ even when $\text{rank}(\mathbf{Y}) = d$. Nonetheless, we will now verify that a non-linear k_y introduces attraction-repulsion characteristics like those in ARDR methods.

⁶To be precise, the kernel matrix is $\mathbf{K}_Y = \mathbf{C}(\mathbf{G}_Y)\mathbf{C}$.

5.1 PCA with Two Kernels

Recall the PCA objective function $\mathcal{L}_{PCA}(\mathbf{X}, \mathbf{Y}) = \|\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\|_F^2$. We now replace the centered Gram matrices $\mathbf{C}\mathbf{G}_X\mathbf{C}$ and $\mathbf{C}\mathbf{G}_Y\mathbf{C}$ by their kernel⁷ counterparts \mathbf{K}_X and \mathbf{K}_Y . Our loss function is then $\mathcal{L}_{PCA}^{2-Kernel}(\mathbf{X}, \mathbf{Y}) = \|\mathbf{K}_X - \mathbf{K}_Y\|_F^2$, leading to a gradient of the form

$$\nabla_{y_i} = - \sum_j (k_x(x_i, x_j) - k_y(y_i, y_j)) \cdot \frac{d k_y(\|y_i - y_j\|_2^2)}{d \|y_i - y_j\|_2^2} \cdot (y_i - y_j) \quad (7)$$

It was shown in [52] that applying these gradients with the UMAP optimization scheme provides embeddings that are indistinguishable from the standard UMAP embeddings. However, we show in Figure 3 that this does not hold when optimizing all $O(n^2)$ pairwise forces. This is in line with the observations made in [48] regarding what loss UMAP is actually optimizing – applying some of the gradients in Eq. 7 gives embeddings indistinguishable from UMAP but optimizing all n^2 pairs of points does not. Thus, reproducing ARDR embeddings in terms of classical techniques requires a method that prioritizes local/global relationships similarly to tSNE and UMAP.

5.2 LLE with Two Kernels

LLE is therefore a natural choice: it finds the \mathbf{Y} that preserves the local relationships in \mathbf{X} . We now present the gradients that optimize the LLE objective under two kernels.

For simplicity’s sake, we will assume⁸ that the constraint $\frac{1}{n}\mathbf{Y}^\top\mathbf{Y} = \mathbf{I}$ applies in kernel space. The closest suitable constraint is then $\frac{1}{n}\mathbf{K}_Y = \mathbf{I}$, as this implies that $\psi(\mathbf{Y})$ is an orthogonal basis and will also have covariance matrix \mathbf{I} . Notice that the diagonal of both \mathbf{K}_Y and \mathbf{I} is always 1, implying that the constraint requires the off-diagonals of \mathbf{K}_Y to be 0. Thus, we will approximate the solution to LLE with two kernels⁹ by minimizing the loss $\mathcal{L}_{LLE}(\mathbf{W}, \mathbf{Y}) = \text{Tr}(\mathbf{M}\mathbf{K}_Y) + \frac{1}{n} \sum_{i,j} k_y(\|y_i - y_j\|_2^2)$, for $\mathbf{M} = (\mathbf{I} - \mathbf{W})^\top(\mathbf{I} - \mathbf{W})$.

Given this loss function, assume that we have the \mathbf{W} that represents nearest neighborhoods in \mathbf{X} . We now want to use gradient descent to find the \mathbf{Y} such that $\mathcal{L}_{LLE}(\mathbf{W}, \mathbf{Y})$ is minimized. Our next lemma shows that this has a very natural interpretation in the A/R framework.

Lemma 4. Let \mathbf{Y} and \mathbf{M} be defined as above, let $\mathbf{V} = \mathbf{W}^\top\mathbf{W}$, and let c be a constant. Then the gradient of $\text{Tr}(\mathbf{M}\mathbf{K}_Y) + \frac{1}{n} \sum_{i,j} k_y(\|y_i - y_j\|_2^2)$ with respect to y_i is given by

$$\nabla_{y_i} = c \sum_j \left(w_{ij} + w_{ji} - v_{ij} - \frac{1}{n} \right) \frac{d k_y(\|y_i - y_j\|_2^2)}{d \|y_i - y_j\|_2^2} (y_i - y_j).$$

The proof can be found in Section B.2 of the supplementary material.

⁷Note that $\mathbf{C}\mathbf{A}\mathbf{C}$ is still a psd matrix for psd \mathbf{A} ; thus \mathbf{K}_X and \mathbf{K}_Y can implicitly be double-centered.

⁸It is unclear to us if the constraint should apply in kernel space $\psi(\mathbf{Y})$ or in the embedding space \mathbf{Y} .

⁹We recognize that this is a simplistic view of the problem. Nonetheless, even this simple version proves sufficient for emulating ARDR gradients.

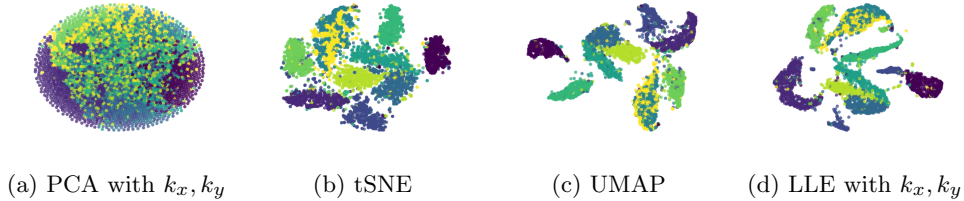


Fig. 3: Left: Embedding by grad. descent of the PCA objective function with the UMAP k_x and k_y functions. Middle: tSNE and UMAP embeddings. Right: Embedding by grad. descent of the LLE objective function with the UMAP k_x and k_y functions. The dataset is 5K randomly chosen points from the MNIST dataset. We observe similar separation of the classes in LLE with two kernels vs. tSNE and UMAP.

We take a moment to consider the striking similarities between this and UMAP’s gradients. First, note that for $k_y \sim (1 - \|y_i - y_j\|_2^2)^{-1}$ as in UMAP, the derivative with respect to the distance incurs a minus sign. Given this, the terms w_{ij} and w_{ji} constitute our attraction scalars, i.e. the attraction between y_i and y_j depends *linearly* on how much x_j (resp. x_i) contributes to x_i ’s (resp. x_j ’s) local neighborhood under k_x . Indeed, this is precisely what the UMAP attraction scalars constitute. Furthermore, in UMAP’s [implementation](#)¹⁰, when y_i is attracted to y_j , y_j is also attracted to y_i [52].

Now consider the repulsive terms in Lemma 4 given by $v_{ij} + 1/n = [\mathbf{W}^\top \mathbf{W}]_{ij} + 1/n$. Since \mathbf{W} is an adjacency matrix with k entries in each row, the first term is equal to the sum of the weights over all paths of length 2 from x_i to x_j . That is, v_{ij} imposes repels y_i from y_j based on how densely connected x_i and x_j are. The second repulsion scalar $1/n$ is simply an average repulsion from all points in \mathbf{Y} . Now recall that UMAP’s repulsion heuristic samples a constant number of points from \mathbf{Y} for each y_i . For distant points, these repulsions are effectively 0, implying that UMAP’s practical repulsions are, in expectation, somewhere between the v_{ij} and $1/n$ terms from Lemma 4.

We visualize this in similarity Figure 3, where we show the tSNE, UMAP and 2-kernel PCA and LLE embeddings on 5K points from the MNIST training set. For the 2-kernel implementations, we simply calculate the loss function the corresponding loss function in Pytorch and run standard gradient descent on it. Importantly, we use *none* of the heuristics that are present in the tSNE and UMAP algorithms [2, 52, 67]. The message is that the most vanilla implementation of LLE with UMAP’s kernels adequately reproduces ARDR embeddings.

6 Conclusion and Future Work

Before discussing our conjectures and future work, we mention that they are directed at our primary open question: “What information does an optimal ARDR-like embedding give regarding its input?” In order to answer this, one must have a definition of what it means to find an ‘optimal ARDR-like’ embedding. To that end, we point out that there seems to be a fundamental divide between the modern and classical

¹⁰This is not theoretically motivated but seems to provide better convergence.

DR methods and that quantifying how ‘similar’ two algorithms’ embeddings are is a difficult task [52, 53]. Thus, in the absence of consensus on a metric that can do this, we will discuss our conjectures using the theoretical metric f^* that optimally differentiates between classical and modern DR methods. Although it is infeasible to find such a metric in practice, identifying a measure that adequately approximates f^* should serve to make our conjectures falsifiable. We note that finding a metric that resembles f^* is a major open question and has been discussed in [49, 53, 70, 78].

6.1 Towards an Optimal Metric

Let \mathbf{Y}_1 and \mathbf{Y}_2 be two embeddings obtained from algorithms \mathcal{A}_1 and \mathcal{A}_2 on the same input, i.e. $\mathcal{A}_1(\mathbf{X}) = \mathbf{Y}_1$ and $\mathcal{A}_2(\mathbf{X}) = \mathbf{Y}_2$. We now define the space of metrics¹¹ \mathcal{M} that accept \mathbf{Y}_1 and \mathbf{Y}_2 and output a value in $[0, 1]$ ¹². We now consider the optimal metric $f^* \in \mathcal{M}$ as the one that maximally differentiates between classical DR methods (PCA, LLE, ISOMAP, LE, MDS, etc.) and the modern ones (tSNE, UMAP, etc.). That is, in expectation over the set of all inputs \mathcal{X} , f^* gives maximal scores when the embeddings come from the same class of DR methods and minimal scores when they come from different classes of methods¹³. In some sense, f^* is a measure for the fundamental difference in embeddings between modern and classical DR methods. Equipped with this definition, we now proceed to our conjectures regarding the essence underlying tSNE and UMAP.

6.2 Conjectures

In Section 5, we showed that pairing the classical DR methods with a kernel on \mathbf{Y} gives gradients that have a very similar form to those in ARDR methods. Furthermore, we verified that minimizing the objective of LLE with UMAP’s k_x, k_y -kernels gives similar embeddings as UMAP and tSNE. This implies that the primary difference between the classical and ARDR methods is the kernel on \mathbf{Y} . Specifically, let f^* be as defined in Section 6.1. Then,

Conjecture 1. *Let $\mathbb{E}_{x \in \mathcal{X}} [\nabla_{y_i}^{umap}]$ be the expected gradient calculated during an epoch of UMAP for a given embedding \mathbf{Y} . Let $\nabla_{y_i}^{lle}$ be defined as in Lemma 4. Then $\mathbb{E}_{x \in \mathcal{X}} [\langle \nabla_{y_i}^{umap}, \nabla_{y_i}^{lle} \rangle_F] > 0$, where $\langle \cdot, \cdot \rangle_F$ denotes the Frobenius inner product. Furthermore, embeddings found by gradient descent on LLE with the UMAP kernels will, under f^* , have high similarity to embeddings found by UMAP and low similarity to those found by classical methods.*

¹¹We assume that these metrics have some standard properties. They should be (1) symmetric to the inputs, i.e. $f(\mathbf{Y}_1, \mathbf{Y}_2) = f(\mathbf{Y}_2, \mathbf{Y}_1)$; (2) invariant to the magnitude of the embeddings, i.e. $f(\mathbf{Y}_1, \mathbf{Y}_2) = f(a\mathbf{Y}_1, a\mathbf{Y}_2)$ for $a \in \mathbb{R}$; (3) invariant to orientation, i.e. $f(\mathbf{Y}_1, \mathbf{Y}_2) = f(\mathbf{O}\mathbf{Y}_1, \mathbf{O}\mathbf{Y}_2)$ for any orthonormal transformation \mathbf{O} .

¹²A score of 0 implies that \mathbf{Y}_1 is maximally dissimilar from \mathbf{Y}_2 and 1 means that $\mathbf{Y}_1 = \mathbf{Y}_2$.

¹³

$$f^* = \arg \sup_{f \in \mathcal{M}} \mathbb{E}_{\substack{x \in \mathcal{X} \\ \mathcal{A}_c, \mathcal{A}'_c \in \mathcal{A}_{classic} \\ \mathcal{A}_m, \mathcal{A}'_m \in \mathcal{A}_{ardr}}} [f(\mathcal{A}_c(x), \mathcal{A}'_c(x)) + f(\mathcal{A}_m(x), \mathcal{A}'_m(x)) - 2f(\mathcal{A}_c(x), \mathcal{A}_m(x))] \quad (8)$$

That is, we claim that a gradient step for LLE with the UMAP kernels is, on average, going to act on the embedding similarly to UMAP and that the final embeddings will have similar structure under f^* . This is a formalization of the statement ‘UMAP approximates LLE with two kernels’.

Although the above conjecture describes how one could frame UMAP in terms of the classical algorithms, it states nothing about convergence guarantees. Indeed, we have seen that performing PCA in the ARDR framework with gradient descent converges to the true PCA embedding. It remains to be seen if this holds in the presence of non-linear k_y kernels. However, if Conjecture 1 is true, then we believe that there should exist a direct method similar to LLE that emulates tSNE/UMAP:

Conjecture 2. *Let \mathcal{A} be an ARDR algorithm that has high similarity to UMAP and low similarity to classical methods under f^* and let \mathcal{L} be \mathcal{A} ’s effective loss function. Let OPT_x be the embedding with minimal cost for \mathcal{A} on $x \in \mathcal{X}$ under \mathcal{L} . Then there exists an algorithm \mathcal{A}' that returns $\mathcal{A}'(x) = \mathbf{Y}'_x$ without iterative optimizations such that $\mathcal{L}(\mathbf{Y}'_x) \in (1 \pm \varepsilon)\mathcal{L}(OPT_x)$ for all $x \in \mathcal{X}$.*

By *effective* loss function, we mean the loss that is being optimized by the specific heuristics and implementation of \mathcal{A} ; we refer to [48] for an example regarding UMAP. This conjecture is a formalization of the statement ‘an algorithm similar to UMAP can be performed without gradient descent and can provably approximate the global minimum of UMAP’.

If we assume the above conjectures are true, we now propose that we can transfer the explainability of PCA and LLE to the ARDR setting. That is, if one can get the same quality embedding by solving for it directly, then one would understand how well the embedding actually preserves the structure of the dataset. In a formal sense, we believe that the major open question surrounding tSNE and UMAP is the following:

Question 1. *Let \mathcal{A} be an ARDR algorithm that is similar to tSNE/UMAP under f^* and let $OPT_{\mathcal{A}}(x)$ be the optimal embedding under \mathcal{A} for input x . Now consider all $\{x_1, x_2, \dots, x_n\} \in \mathcal{X}$ such that $OPT_{\mathcal{A}}(x_1) = OPT_{\mathcal{A}}(x_2) = \dots = OPT_{\mathcal{A}}(x_n)$. Then what characteristics must be consistent across all $x \in \{x_1, \dots, x_n\}$?*

This is perhaps the most fundamental explainability question that one can ask: “given the output of the explainability method, what can one say about the input?”

We note that our conjectures directly help to answer this question. First, our discussion on metrics describes what characteristics a ‘similar’ algorithm to tSNE/UMAP must exhibit. Second, we believe that LLE with two kernels is precisely the ARDR algorithm that is ‘similar’ to tSNE/UMAP. Lastly, we conjecture that such an ARDR algorithm can be solved directly, i.e. without relying on attractions and repulsions. This would allow one to formally state which inputs map to the same embedding and, inevitably, what qualities must be consistent across all of these inputs.

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A PCA theoretical results

A.1 Proof of Theorem 1

Proof. The proof relies on a change of basis in the first d components of \mathbf{CX} and \mathbf{CY} . Let $\mathbf{CX} = \mathbf{U}_X \boldsymbol{\Sigma}_X \mathbf{V}_X^\top$ and $\mathbf{CY} = \mathbf{U}_Y \boldsymbol{\Sigma}_Y \mathbf{V}_Y^\top$ be the SVD of \mathbf{CX} and \mathbf{CY} . Then $\mathbf{CG}_X \mathbf{C} = \mathbf{U}_X \boldsymbol{\Sigma}_X^2 \mathbf{U}_X^\top$ and $\mathbf{CG}_Y \mathbf{C} = \mathbf{U}_Y \boldsymbol{\Sigma}_Y^2 \mathbf{U}_Y^\top$.

We now decompose $\mathbf{CX} = \mathbf{U}_X^+ \boldsymbol{\Sigma}_X^+ (\mathbf{V}_X^+)^{\top} + \mathbf{U}_X^- \boldsymbol{\Sigma}_X^- (\mathbf{V}_X^-)^{\top}$, where the $+$ superscript corresponds to the first d components of \mathbf{CX} and the $-$ superscript corresponds to the final $D-d$ components. Note that now \mathbf{U}_X^+ is an orthogonal transformation in the same subspace as \mathbf{U}_Y . This allows us to define $\mathbf{O} = \mathbf{U}_Y (\mathbf{U}_X^+)^{\top}$ as the change of basis from $\mathbf{CG}_X \mathbf{C}$ to $\mathbf{CG}_Y \mathbf{C}$, providing the following characterization of $L_{PCA}(\mathbf{X}, \mathbf{Y})$:

$$L_{PCA}(\mathbf{X}, \mathbf{Y}; \mathbf{O}) = \|\mathbf{OCG}_X \mathbf{CO}^\top - \mathbf{CG}_Y \mathbf{C}\|_F^2 = \|(\mathbf{U}_Y ((\boldsymbol{\Sigma}_X^+)^2 - \boldsymbol{\Sigma}_Y^2) \mathbf{U}_Y^\top) + f(\mathbf{X})\|_F^2 \quad (9)$$

where $f(\mathbf{X})$ is independent of \mathbf{Y} . The minimum of Equation 9 over \mathbf{Y} occurs when \mathbf{CY} has the same singular values as \mathbf{OCX} in the first d components, which is exactly the PCA projection of \mathbf{X} up to orthonormal transformation.

It remains to show that PCA is invariant to orthonormal transformations. Let $\mathbf{X}' = \mathbf{OX}$. Then $\mathbf{G}'_X = \mathbf{X}'(\mathbf{X}')^\top = \mathbf{OG}_X \mathbf{O}^\top = \mathbf{G}_X$, since Gram matrices are unique up to orthogonal transformations. \square

A.2 PCA gradient derivation

Let \mathbf{Y} be *any* set of points in $\mathbb{R}^{n \times d}$. Then the gradient with respect to \mathbf{Y} of $\mathcal{L}_{PCA}(\mathbf{X}, \mathbf{Y}) = \|\mathbf{CG}_X \mathbf{C} - \mathbf{CG}_Y \mathbf{C}\|_F^2$ is obtained by

$$\begin{aligned} d \mathcal{L}_{PCA}(\mathbf{X}, \mathbf{Y}) &= d \|\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\|_F^2 \\ &= d \left((\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C})^\top : (\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}) \right) \\ &= 2(\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C})^\top : d(\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}) \\ &= -2\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C} : d\mathbf{G}_Y \\ &= -4\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{CY} : d\mathbf{Y} \\ \frac{d \mathcal{L}_{PCA}(\mathbf{X}, \mathbf{Y})}{d\mathbf{Y}} &= -4\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{CY} \end{aligned}$$

where $:$ represents the Frobenius inner product and the centering matrices cancel due to idempotence.

A.3 Proof of Lemma 2

Proof. We use the well-known fact that $\mathbf{CG}_Y \mathbf{C} = -\frac{1}{2}\mathbf{CD}_Y \mathbf{C}$, for squared Euclidean distance matrix \mathbf{D}_Y , to express the gradient in terms of $(y_i - y_j)$ vectors.

$$\nabla = \alpha \mathbf{L} : d \left(-\frac{1}{2}\mathbf{CD}_Y \mathbf{C} \right)$$

$$\begin{aligned}
&= -\frac{\alpha}{2} \mathbf{L} : d\mathbf{D}_Y && (\mathbf{C} \text{ cancels by idempotence}) \\
\implies \nabla_{y_i} &= -\alpha \sum_j [\mathbf{L}]_{ij} (y_i - y_j)
\end{aligned}$$

□

A.4 Proof of Theorem 1

We start by confirming that the PCA embedding is a minimum of the gradient. Recall that our gradient is of the form

$$\nabla_{\mathbf{Y}} = 4\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\mathbf{Y}.$$

If $\mathbf{C}\mathbf{Y}$ is the PCA embedding of $\mathbf{C}\mathbf{X}$, then $\mathbf{C}\mathbf{Y}$'s d singular values should be $\mathbf{C}\mathbf{X}$'s top- d singular values. Given this, $\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}$ will only have singular values corresponding to the remaining $D - d$ diagonal positions. However, notice that \mathbf{Y} 's singular values are still in the first d coordinates. Since \mathbf{G}_X and \mathbf{G}_Y have the same orthonormal basis, we see that $\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}$ is orthogonal to \mathbf{Y} , giving us 0 gradient when \mathbf{Y} is the PCA embedding.

Next, we show that any minimum of the objective function must be a global minimum. This is a simple extension of the proof of lemma 1. Assume for the sake of contradiction that \mathbf{Z} is a global minimum of \mathcal{L}_{PCA} that is *not* equal to the PCA embedding. Then $\mathbf{Z} \neq \mathbf{O}\mathbf{Y}$ for some orthonormal transformation \mathbf{O} , implying that the singular values of \mathbf{Z} do not equal those of \mathbf{X} in the first d components. Thus, there is a \mathbf{Z}' in the ε -ball of \mathbf{Z} that better approximates the singular values of \mathbf{X} . Thus, \mathbf{Z} cannot be a minimum.

Given this, we now want to show that there always exists a learning rate value such that a step of gradient descent will decrease our cost. Let $\mathbf{Y}' = (\mathbf{I} + \gamma\mathbf{L})\mathbf{Y}$ be the embedding after one step of gradient descent with learning rate γ and $\mathbf{L} = \mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}$.

$$\begin{aligned}
\mathcal{L}_{PCA}(\mathbf{X}, \mathbf{Y}') < \mathcal{L}_{PCA}(\mathbf{X}, \mathbf{Y}) &\rightarrow \|\mathbf{C}(\mathbf{G}_X - \mathbf{Y}'\mathbf{Y}'^\top)\mathbf{C}\|_F^2 < \|\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\|_F^2 \\
\|\mathbf{C}(\mathbf{X}\mathbf{X}^\top - \mathbf{Y}'\mathbf{Y}'^\top)\mathbf{C}\|_F^2 &= \|\mathbf{C}(\mathbf{G}_X - ((\mathbf{I} + \gamma\mathbf{L})\mathbf{Y})(\mathbf{I} + \gamma\mathbf{L})\mathbf{Y}^\top)\mathbf{C}\|_F^2 \\
&= \|\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C} - \gamma\mathbf{C}(\mathbf{L}\mathbf{G}_Y - \mathbf{G}_Y\mathbf{L}^\top - \gamma\mathbf{L}\mathbf{G}_Y\mathbf{L}^\top)\mathbf{C}\|_F^2
\end{aligned}$$

This expression subtracts $\gamma\mathbf{C}(\mathbf{L}\mathbf{G}_Y + \mathbf{G}_Y\mathbf{L}^\top + \gamma\mathbf{L}\mathbf{G}_Y\mathbf{L}^\top)\mathbf{C}$ from our initial matrix $\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}$. Thus, the overall Frobenius norm of the difference will be smaller than that of $\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}$ if the two matrices point in the same direction. Said otherwise, we want to find a γ such that

$$0 < \langle \mathbf{C}(\mathbf{L}\mathbf{G}_Y + \mathbf{G}_Y\mathbf{L}^\top + \gamma\mathbf{L}\mathbf{G}_Y\mathbf{L}^\top)\mathbf{C}, \mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C} \rangle_F$$

Applying the properties of the trace as the Frobenius inner product, we get

$$\begin{aligned}
& \text{Tr} \left[(\mathbf{C}(\mathbf{L}\mathbf{G}_Y + \mathbf{G}_Y\mathbf{L}^\top + \gamma\mathbf{L}\mathbf{G}_Y\mathbf{L}^\top)\mathbf{C})^\top \mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C} \right] = \\
& = \text{Tr} \left[(\mathbf{C}(\mathbf{L}\mathbf{G}_Y + \mathbf{G}_Y\mathbf{L}^\top + \gamma\mathbf{L}\mathbf{G}_Y\mathbf{L}^\top)\mathbf{C})^\top \mathbf{L} \right] \\
& = \text{Tr} \left[2\mathbf{L}^2\mathbf{G}_Y\mathbf{L} + \gamma\mathbf{L}^2\mathbf{G}_Y\mathbf{L} \right] \\
& = \text{Tr} \left[2\mathbf{L}^2\mathbf{G}_Y + \gamma\mathbf{L}^3\mathbf{G}_Y \right] \\
& = \text{Tr} \left[2(\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C})^2\mathbf{G}_Y \right] + \gamma\text{Tr} \left[(\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C})^3\mathbf{G}_Y \right] \tag{10}
\end{aligned}$$

where we cancelled the \mathbf{C} matrices by idempotence with respect to the \mathbf{C} 's in \mathbf{L} .

Now notice that the first term is necessarily non-negative as it is the trace of a positive semidefinite matrix. Furthermore, it will only be zero if $\mathbf{G}_X = \mathbf{G}_Y$ or $\mathbf{G}_Y = 0$. In both of these cases our gradient descent has completed, so we can assume the first term to be strictly positive.

If $\mathbf{G}_Y \leq \mathbf{G}_X$, then the second is also positive semidefinite and non-negative. This is intuitively clear, as $\mathbf{G}_Y \leq \mathbf{G}_X$ implies that our problem is entirely convex. Indeed, this means that as long as our learning rate isn't so large that it 'overshoots' the convex subspace, we are guaranteed to minimize our loss when $\mathbf{G}_Y \leq \mathbf{G}_X$. In the alternative setting where $\mathbf{G}_Y \not\leq \mathbf{G}_X$, we simply need to choose a small enough γ such that expression 10 remains positive. This concludes the proof.

A.5 Proof of Lemma 3

The proof relies on defining $\mathbf{G}'_X = \mathbf{G}_X + \mathbf{R}_X$ and $\mathbf{G}'_Y = \mathbf{G}_Y + \mathbf{R}_Y$ as the sum of the original Gram matrices plus residual matrices \mathbf{R} . Then

$$\|\mathbf{R}_X\|_F^2 = (1 + \epsilon)\|\mathbf{G}_X - \mathbf{G}_X^k\|_F^2 = (1 + \epsilon) \sum_{i=k+1}^n \sigma_{xi}^2$$

where σ_{xi} is the i -th singular value of \mathbf{G}_X . Notice that since $\mathbf{G}_Y^k = \mathbf{G}_Y$ for $k \leq d$, we have $\mathbf{R}_Y = 0$.

Plugging these in, we get

$$\begin{aligned}
\langle \nabla, \nabla' \rangle_F &= \text{Tr} \left[\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\mathbf{Y}\mathbf{Y}^\top\mathbf{C}(\mathbf{G}'_X - \mathbf{G}'_Y)\mathbf{C} \right] \\
&= \text{Tr} \left[\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\mathbf{G}_Y\mathbf{C} \left[(\mathbf{G}_X - \mathbf{G}_Y) - (\mathbf{R}_X - \mathbf{R}_Y) \right] \mathbf{C} \right] \\
&= \text{Tr} \left[(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\mathbf{G}_Y\mathbf{C} \left[(\mathbf{G}_X - \mathbf{G}_Y) - \mathbf{R}_X \right] \mathbf{C} \right] \\
&= \text{Tr} \left[((\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\mathbf{G}_Y\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)) - ((\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\mathbf{G}_Y\mathbf{C}\mathbf{R}_X) \mathbf{C} \right] \\
&= \text{Tr} \left[(\mathbf{G}_X - \mathbf{G}_Y)^2\mathbf{C}\mathbf{G}_Y\mathbf{C} \right] - \text{Tr} \left[\mathbf{C}(\mathbf{G}_X - \mathbf{G}_Y)\mathbf{C}\mathbf{G}_Y\mathbf{C}\mathbf{R}_X \right]
\end{aligned}$$

where we perform rearrangements and cancel one of the \mathbf{C} 's due to the trace being invariant to cyclic permutations.

Notice that the first term $\text{Tr}((G_X - G_Y)^2 C G_Y C)$ must be positive as it is the trace of a positive semi-definite matrix. It then suffices to show that

$$\text{Tr}((G_X - G_Y)^2 C G_Y C) > |\text{Tr}(C(G_X - G_Y) C G_Y C R_X)|$$

If this condition is satisfied, then we have that $\langle \nabla, \nabla' \rangle_F > 0$, allowing us to employ subgradient descent methods.

$$\begin{aligned} \langle \nabla, \nabla' \rangle &\geq \text{Tr}((G_X - G_Y)^2 C G_Y C) - |\text{Tr}(C(G_X - G_Y) C G_Y C R_X)| \\ &\geq \text{Tr}((G_X - G_Y)^2 C G_Y C) - c \cdot \|(G_X - G_Y)(C G_Y C)^{1/2}\|_F \cdot \|(C G_Y C)^{1/2} R_X\|_F \\ &= \|(G_X - G_Y)(C G_Y C)^{1/2}\|_F^2 - c \|(G_X - G_Y)(C G_Y C)^{1/2}\|_F \cdot \|(C G_Y C)^{1/2} R_X\|_F \\ &= \|(G_X - G_Y)(C G_Y C)^{1/2}\|_F \left(\|(C G_Y C)^{1/2} (G_X - G_Y)\|_F - c \cdot \|(C G_Y C)^{1/2} R_X\|_F \right) \end{aligned}$$

where $c = \|C\|_F^2$.

Since the first term $\|(G_X - G_Y)(C G_Y C)^{1/2}\|_F$ is necessarily positive, we have that $\langle \nabla, \nabla' \rangle_F \geq 0$ as long as $\|(C G_Y C)^{1/2} (G_X - G_Y)\|_F - c \cdot \|(C G_Y C)^{1/2} R_X\|_F \geq 0$. We can also remove the c scalar, since we know that $c > 1$. This gives us the necessary condition

$$\|(C G_Y C)^{1/2} (G_X - G_Y)\|_F - \|(C G_Y C)^{1/2} R_X\|_F \geq 0 \implies \langle \nabla, \nabla' \rangle_F \geq 0 \quad (11)$$

We can think of $(C G_Y C)^{1/2}$ as any dataset with the same principal components as CY . So this necessary condition is effectively saying that the inner product between CY and $(G_X - G_Y)$ must be bigger than the inner product between CY and R_X . This intuitively makes sense. Consider that $G_X - G_Y$ is the amount of error in our current projection CY . Meanwhile, R_X is an ϵ -approximation of G_X^k , the optimal low-rank representation of G_X . So our necessary condition states that as long as G_Y is not an ϵ -approximation of G_X , we can continue to use the sublinear-time approximation of G_X to approximate the gradient ∇ .

Said otherwise, if G_Y is sufficiently different from G_X , then $\langle \nabla, \nabla' \rangle_F$ is positive. If not, then we have that G_Y approximates G_X^k , the optimal low-rank approximation of G_X . We formalize this below.

Let G_Y be such that $\|G_X - G_Y\|_F^2 \geq (1 + \alpha) \|G_X - G_X^k\|_F^2$ for $\alpha > 0$. Then we want to solve for the α that makes equation 11 necessarily positive. This is equivalent to finding the α that satisfies $\min \|(G_X - G_Y)(C G_Y C)^{1/2}\|_F = \max \|(C G_X C)^{1/2} R_X\|_F$. We then obtain a lower bound for the minimum:

$$\begin{aligned} \|(G_X - G_Y)(C G_Y C)^{1/2}\|_F &\geq \sigma_{\min} \left((C G_Y C)^{1/2} \right) \cdot \|G_X - G_Y\|_F \\ &\geq \sigma_{\min} \left((C G_Y C)^{1/2} \right) \cdot \sqrt{(1 + \alpha) \|G_X - G_X^k\|_F^2} \end{aligned}$$

and an upper bound for the maximum:

$$\begin{aligned}
\|(CG_Y C)^{1/2} R_X\|_F &\leq \|(CG_Y C)^{1/2}\|_2 \|R_X\|_F \\
&\leq \|(CG_Y C)^{1/2}\|_2 \cdot \sqrt{(1+\epsilon) \|G_X - G_X^k\|_F^2} \\
&\leq \sigma_{max} \left((CG_Y C)^{1/2} \right) \cdot \sqrt{(1+\epsilon) \|G_X - G_X^k\|_F^2}
\end{aligned}$$

Setting the lower bound equal to the upper bound and solving for α tells us that equation 11 is greater than 0 when α satisfies

$$\begin{aligned}
\alpha &> \frac{\lambda_{y_1}}{\lambda_{y_k}} \cdot \frac{(1+\epsilon) \|G_X - G_X^k\|_2^2}{\|G_X - G_X^k\|_F^2} - 1 \\
&= (1+\epsilon) \frac{\lambda_{y_1}}{\lambda_{y_k}} - 1
\end{aligned}$$

This means that our gradient ∇' will be in line with the true gradient ∇ as long as G_Y satisfies the following condition:

$$\|G_X - G_Y\|_F^2 \geq (1+\epsilon) \frac{\lambda_{y_1}}{\lambda_{y_k}} \|G_X - G_X^k\|_F^2$$

As long as this is true, the gradient will push Y into the direction of the PCA embedding of X in k dimensions. If we initialize our Y such that $\lambda_{y_1} \approx \lambda_{y_k}$, then we know they will slowly diverge over the course of gradient descent. Thus, we can upper bound the ratio $\lambda_{y_1}/\lambda_{y_k} \leq \lambda_{x_1}/\lambda_{x_k}$. This gives us our final condition for convergence:

$$\langle \nabla, \nabla' \rangle_F \text{ will be greater than 0 as long as } \|G_X - G_Y\|_F^2 \geq (1+\epsilon) \frac{\lambda_{x_1}}{\lambda_{x_k}} \|G_X - G_X^k\|_F^2 \quad (12)$$

B LLE Theoretical Results

B.1 LLE Derivation

If $x_i \approx \sum_k w_{ik} x_{ik}$ is the representation of x_i by a linear combination of its nearest neighbors, then we want to find the \mathbf{Y} such that $y_i \approx \sum_k w_{ik} y_{ik}$. Treating this as an optimization problem, we can write

$$\min \|\mathbf{Y} - \mathbf{WY}\|_F^2 \quad \text{s.t.} \quad \mathbf{Y}^\top \mathbf{Y} = \mathbf{I}.$$

By applying a Lagrangian to the constraint and taking the gradient, we have

$$\begin{aligned}
\min \text{Tr} \left((\mathbf{I} - \mathbf{W}) \mathbf{G}_Y (\mathbf{I} - \mathbf{W})^\top \right) + \text{Tr} \left(\mathbf{\Lambda} (\mathbf{I} - \mathbf{Y}^\top \mathbf{Y}) \right) \\
\Rightarrow \nabla_{\mathbf{Y}} = 2\mathbf{MY} - 2\mathbf{Y}^\top \mathbf{\Lambda}
\end{aligned}$$

$$\text{set } \nabla_{\mathbf{Y}} \text{ to } 0 \implies \mathbf{M}\mathbf{Y} = \mathbf{Y}^\top \mathbf{\Lambda}$$

where $\mathbf{M} = (\mathbf{I} - \mathbf{W})^\top (\mathbf{I} - \mathbf{W})$. Since we are minimizing the objective, the embedding \mathbf{Y} is given by the eigenvectors of $\mathbf{M} = (\mathbf{I} - \mathbf{W})^\top (\mathbf{I} - \mathbf{W})$ that correspond to the smallest d positive eigenvalues as these represent the smallest Lagrangians. Note that $(\mathbf{I} - \mathbf{W})$ is a graph Laplacian matrix and therefore \mathbf{M} has at least 1 zero eigenvalue.

B.2 Proof of Lemma 4

Proof. We seek the gradient of

$$\mathcal{L}_{LLE}(\mathbf{Y}) = \text{Tr}(\mathbf{M}\mathbf{K}_Y) + \frac{1}{n} \text{Tr}((\mathbf{I} - \mathbf{K}_Y)^2)$$

We start with the first term. By differentiating the Frobenius inner product, we have

$$\frac{\partial \text{Tr}(\mathbf{M}\mathbf{K}_Y)}{\partial \mathbf{Y}} = \mathbf{M} : \frac{\partial \mathbf{K}_Y}{\partial \mathbf{Y}}. \quad (13)$$

Now consider that the (i, j) -th entry of \mathbf{K}_Y is a function of $\|y_i - y_j\|_2^2$ and therefore only depends on the (i, j) -th entry of \mathbf{D}_Y . Thus,

$$\frac{\partial \mathbf{K}_Y}{\partial \mathbf{Y}} = \frac{\partial \mathbf{K}_Y}{\partial \mathbf{D}_Y} \odot \frac{\partial \mathbf{D}_Y}{\partial \mathbf{Y}},$$

where \odot is the Hadamard element-wise matrix product. We plug this into Eq. 13 and rearrange terms to get

$$\frac{\partial \text{Tr}(\mathbf{M}\mathbf{K}_Y)}{\partial \mathbf{Y}} = \left[\frac{\partial \mathbf{K}_Y}{\partial \mathbf{D}_Y} \odot \mathbf{M} \right] : \frac{\partial \mathbf{D}_Y}{\partial \mathbf{Y}}.$$

We now use the intuition developed in Section 3.3.1 to point out that this corresponds to the gradient acting on point y_i as

$$\nabla_{y_i}(\text{Tr}(\mathbf{M}\mathbf{K}_Y)) = c \sum_j m_{ij} \frac{d k_y(\|y_i - y_j\|_2^2)}{d \|y_i - y_j\|_2^2} (y_i - y_j) \quad (14)$$

Now recall that $\mathbf{M} = (\mathbf{I} - \mathbf{W})^\top (\mathbf{I} - \mathbf{W}) = -\mathbf{W} - \mathbf{W}^\top + \mathbf{I} + \mathbf{W}^\top \mathbf{W}$. Thus, we can write Eq. 14 as

$$\nabla_{y_i}(\text{Tr}(\mathbf{M}\mathbf{K}_Y)) = c \sum_j (-w_{ij} - w_{ji} + \mathbb{1}_{i=j} + [\mathbf{W}^\top \mathbf{W}]_{ij}) \frac{d k_y(\|y_i - y_j\|_2^2)}{d \|y_i - y_j\|_2^2} (y_i - y_j),$$

where $\mathbb{1}_{i=j}$ is 1 if $i = j$ and 0 otherwise. However, notice that if $y_i = y_j$ implies that $y_i - y_j$ is 0. Thus, in the $i = j$ setting the gradient will be 0. Thus, we can cancel the $\mathbb{1}$ term from the sum, giving the desired result.

It remains to show the gradient of the second term $\frac{1}{n} \sum_{i,j} k_y(\|y_i - y_j\|_2^2)$. There, it is a simple re-use of the above:

$$\begin{aligned} \frac{\partial(\frac{1}{n} \sum_{i,j} k_y(\|y_i - y_j\|_2^2))}{\partial \mathbf{Y}} &= \frac{1}{n} \text{sum}_{i,j} \frac{\partial(k_y(\|y_i - y_j\|_2^2))}{\partial \mathbf{Y}} \\ \implies \nabla_{y_i} &= -\frac{1}{n} \sum_j \frac{d k_y(\|y_i - y_j\|_2^2)}{d \|y_i - y_j\|_2^2} (y_i - y_j) \end{aligned}$$

□