ON THE RELATION BETWEEN LINEAR DIFFUSION AND POWER ITERATION

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ABSTRACT

Recently, diffusion models have gained popularity due to their impressive generative abilities. These models learn the implicit distribution given by the training dataset, and sample new data by transforming random noise through the reverse process, which can be thought of as gradual denoising. In this work, we examine the generation process as a "correlation machine", where random noise is repeatedly enhanced in correlation with the implicit given distribution. To this end, we explore the linear case, where the optimal denoiser in the MSE sense is known to be the PCA projection. This enables us to connect the theory of diffusion models to the spiked covariance model, where the dependence of the denoiser on the noise level and the amount of training data can be expressed analytically, in the rank-1 case. In a series of numerical experiments, we extend this result to general low rank data, and show that low frequencies emerge earlier in the generation process, where the denoising basis vectors are more aligned to the true data with a rate depending on their eigenvalues. This model allows us to show that the linear diffusion model converges in mean to the leading eigenvector of the underlying data, similarly to the prevalent power iteration method. Finally, we empirically demonstrate the applicability of our findings beyond the linear case, in the Jacobians of a deep, non-linear denoiser, used in general image generation tasks.

1 INTRODUCTION

Recently, diffusion models have gained much popularity as very successful generative models, showcasing impressive performance in image generation tasks (Dhariwal & Nichol, 2021; Ho et al., 2020; Song & Ermon, 2019; Song et al., 2021c). These models learn the implicit distribution given by the training dataset, and sample new data by transforming random noise inputs through a reverse diffusion process, which can be thought of as gradual denoising. More formally, it has been shown in Kadkhodaie et al. (2024) that learning the underlying distribution is equivalent to optimal denoising at all noise levels.

In order to shed more light onto the mechanism behind the success of diffusion models, in this work we analyze the behavior of denoisers in the context of image generation, where pure noise is gradually processed into a sample from a given (implicit) distribution by gradual denoising. Unlike other works, e.g. Kadkhodaie et al. (2024), we focus on the denoiser(s) throughout the generation process, and not only on the final generated data.

To this end, we suggest the following simple model to illustrate our point. Consider the class of linear denoisers, where the optimal denoiser is given by a PCA projection. To simulate the diffusion generation process, we learn a series of projections onto noisy data at different noise levels, and use them to transform pure noise into samples from the underlying distribution. Given this simple model we can inspect the evolution of eigenvectors spanning gradual projections with decreasing noise levels, as well as the distribution of the generated data samples.

We show that the correlation of the noisy basis eigenvectors with their clean version decays as the noise level increases, with a rate determined by the eigenvalues and the size of the training dataset. In other words, we show that low frequencies, corresponding to large eigenvalues, emerge earlier in the reverse process as was empirically observed in (Ho et al., 2020), and analyze how more training data contributes to generalization (Kadkhodaie et al., 2024). Analytically, this corresponds to the spiked covariance model (Johnstone, 2001), in which we bound this decay for the leading eigenvector (corresponding to the largest eigenvalue).

Next, we demonstrate the applicability of our findings to more general, non-linear deep denoisers. Although the network is not linear, its application can be written as a linear operation of the Jacobian calculated on the input image. We empirically show that the aforementioned decay of eigenvector correlations is prevalent also in in the Jacobians of a deep denoiser, in the final stages of image generation, thus showing the relevance of our analysis in a broader context, and not just in a simplified linear case.

2 Background and Related Work

Since their introduction by Sohl-Dickstein et al. (2015), diffusion models have been vastly used in image generation tasks (Dhariwal & Nichol, 2021; Ho et al., 2020; Song & Ermon, 2019; Song et al., 2021c), more general computer vision tasks (Amit et al., 2021; Baranchuk et al., 2022; Brempong et al., 2022; Cai et al., 2020), and in other domains such as natural language processing (Austin et al., 2021; Hoogeboom et al., 2021; Li et al., 2022; Savinov et al., 2022; Yu et al., 2022) and temporal data modeling (Alcaraz & Strodthoff, 2023; Chen et al., 2021; Kong et al., 2021; Rasul et al., 2021; Tashiro et al., 2021). On top of their practical success, different flavors of training and sampling have risen based on interesting theoretical reasoning, e.g., considering the statistical properties of the intermediate data (Song et al., 2021a; Sohl-Dickstein et al., 2015), or by framing the problem in the form of stochastic differential equations (SDEs) (Karras et al., 2022; Song et al., 2021b;c; Chen et al., 2024) or score based generative models (Song & Ermon, 2019; 2020). In this work, we look at diffusion models in the context of iterative denoising, and focus on the properties of the learned denoiser (Milanfar & Delbracio, 2024).

Recently Kadkhodaie et al. (2024) showed that the learned denoising functions are equivalent to a shrinkage operation in a basis adapted to the underlying image. In this sense the diffusion denoiser is an adaptive filter (Milanfar, 2013; Talebi & Milanfar, 2014; 2016). While they focus on the analysis of the nonlinear denoiser at the point of the final generated data, we are interested in the evolution (adaptation) of the denoiser throughout the generation process, and its dependence on the noise level. To this end, we suggest a simple linear denoising model, presented in Section 3. In this case, the (optimal) denoiser does depend on the underlying image, and its dependence on the noise level can be traced analytically, as we show hereafter.

Due to their phenomenal empirical success, some attempts have been devoted towards providing theory supporting the sample and iteration complexity of diffusion models. The current body of work can be generally parted to attaining iteration complexity bounds assuming approximately accurate scores (Li et al., 2024b;a; Chen et al., 2023b; Huang et al., 2024; Benton et al., 2024), and to assessing the sample complexity to learn the score functions (Chen et al., 2023a; Block et al., 2020; Biroli & Mézard, 2023). Among these works, many assume a low dimensional data distribution (Bortoli, 2022; Li & Yan, 2024; Oko et al., 2023; Chen et al., 2023a; Wang et al., 2024), which is a reasonable assumption in practice (see e.g., Pope et al. (2021)). Yet, it might particularly explain the gap between the current iteration bounds and the much lower complexity apparent in practice (Li & Yan, 2024). In our work, we consider linear models and deduce a linear sample complexity bound associated with learning the score function in Sec. 4 and discuss the tradeoffs of the synthesis conversion rate in Sec. 4.1. The previous works mentioned above mainly develop bounds assuming specific samplers and scaling details, which differ from our setting. In addition, they generally bound the Total Variation distance (under varying assumptions on the target distributions), which is not trivial to translate to the generated covariance matrix that we focus on even in the linear Gaussian case (Devroye et al., 2018). The difference in our setting enables us to connect the theory of diffusion models to a broad body of work concerning the spiked covariance model (Johnstone, 2001), and supports the analysis of denoising diffusion as a correlation machine, which is the main purpose of this paper.

In the setting of Statistical Mechanics, Biroli & Mézard (2023) analyses diffusion models in very large dimensions, focusing on the Curie-Weiss model of ferromagnetism. As an introduction to their work, they also discuss a simple linear score model, in the context of the sample complexity of learning the score function. They focus their discussion on the case of Gaussian data, where the eigenvalues of the covariance matrices can be typically characterized. Unlike their work, we consider data that reside in a low dimensional subspace, with no specific distribution, described in Sec. 4.

Power iteration is a fundamental algorithm for approximating the dominant eigenvalue and eigenvector of a matrix. It relies on iteratively multiplying an initial vector by the matrix, where its convergence rate is proportional to the ratio of the largest and second-largest eigenvalues. The method's simplicity and scalability have made it a cornerstone in various fields, including numerical linear algebra, machine learning, and graph theory. For the ease of reading, we include a formal presentation of the method and

discuss its convergence in Appendix A. In this work, we shall show how a linear denoising chain converges in mean to the celebrated power iteration method.

3 Linear Diffusion - Problem Setup

For our analysis, we define the following simple iterative linear generation model. First, define the standard diffusion model. Let q denote the natural data distribution and let $x_0 \sim q$ be a sample from the natural data ($x \in \mathbb{R}^d$). The forward (diffusion) process is defined (Ho et al., 2020) by

$$q(x_t|x_{t-1}) = \mathcal{N}(\sqrt{1-\beta_t}x_{t-1}, \beta_t \mathbf{I}) \tag{1}$$

for some fixed noise schedule $\{\beta_t\}_{t=1}^T$ and $x_0 \sim q$. It can be shown that

$$q(x_t|x_0) = \mathcal{N}(\sqrt{\bar{\alpha}_t}x_0, (1-\bar{\alpha}_t)\mathbf{I}), \tag{2}$$

where $\alpha_t = 1-\beta_t$ and $\bar{\alpha}_t = \Pi_{s=1}^t \alpha_s$. For our simplified model, consider the process (without scaling),

$$q(x_t|x_{t-1}) = \mathcal{N}(x_{t-1}, \sigma_t^2 \mathbf{I}). \tag{3}$$

This implies that $x_t = x_{t-1} + \epsilon_{\sigma_t}$, where $\epsilon_{\sigma_t} \sim \mathcal{N}(0, \sigma_t^2 \mathbf{I})$ for some fixed noise schedule $\{\sigma_t\}_{t=1}^T$. We discard the scaling to comply with previous analysis of the spiked covariance model (Nadler, 2008) (more details in Section 4). This corresponds to the "Exploding Variance" formulation, used with Langevin dynamics to sample data as a variant of score based diffusion models (Song & Ermon, 2019; Song et al., 2021c; Song & Ermon, 2020). We choose to present the "standard" diffusion models in the setting of denoising diffusion Ho et al. (2020) and not using the score-based approach entirely, as we focus our discussion on the qualities of the denoiser.

The reverse (generation) process is defined using a parameterized distribution model p_θ, generally defined by the Markov process

$$p_\theta(x_{0:T}) = p(x_T)\Pi_{t=1}^T p_\theta(x_{t-1}|x_t), \tag{4}$$

$$p_\theta(x_{t-1}|x_t) \triangleq \mathcal{N}(\mu_\theta(x_t,t), \Sigma_\theta(x_t,t)), \tag{5}$$

where $p(x_T) = \mathcal{N}(0, \mathbf{I})$. By choices of parametrization and loss manipulations (see (Ho et al., 2020)), one generally learns to estimate the error $\epsilon_\theta(x_t,t)$, where

$$\mu_\theta(x_t,t) = \frac{1}{\sqrt{\alpha_t}}\Big(x_t - \frac{\beta_t}{\sqrt{1-\bar{\alpha}_t}}\epsilon_\theta(x_t,t)\Big), \tag{6}$$

$\Sigma_\theta(x_t,t) = e_t^2 \mathbf{I}$, and e_t is a designed schedule (usually chosen to be equal to σ_t). Thus, the reverse process can be expressed as a denoising chain

$$D_t(x_t) = \frac{1}{\sqrt{\alpha_t}}\Big(x_t - \frac{\beta_t}{\sqrt{1-\bar{\alpha}_t}}\epsilon_\theta(x_t,t)\Big) + e_t z, \tag{7}$$

where $z \sim \mathcal{N}(0, \mathbf{I})$ and $z_1 = 0$. This is a stochastic denoiser, which preserves the Markovian property of the forward process. Later versions suggested similar (non Markovian) deterministic denoisers, e.g., DDIM (Song et al., 2021a), or more general stochastic denoiser chains, for a continuous forward model (InDI (Delbracio & Milanfar, 2023)).

In our case, we restrict the denoisers to be a linear function of x_t. Thus, the optimal denoiser (in the ℓ_2 sense) is given by the PCA projection onto the target distribution. For the reverse process, we learn a simple PCA denoiser (projection) based on X_{t-1}, which is the cleaner version of the training set $X = \{x_1, ..., x_n\}$ at time $t-1$. Thus, at each time step we learn

$$D_{PCA}^t(x_t) = D_{PCA}^t(x_t; X_{t-1}) = P_t(x_t; X_0 + E_{\bar{\sigma}_t}), \tag{8}$$

where each column in $E_{\bar{\sigma}_t}$ is distributed by $\mathcal{N}(0, \bar{\sigma}_t^2 \mathbf{I})$ and $\bar{\sigma}_t$ is a function of $\{\sigma_s\}_{s=1}^t$. Our simple denoising procedure is based on the sequential application of $P_t \in \mathbb{R}^{d \times d}$, which is the projection on perturbed principal components with respect to the clean data distribution q. It is a deterministic denoiser given the sampling of training data and noises, which does not depend neither on x_t nor on x_0. Nevertheless, this model is relevant in differentiable environments of more complex settings such as DNN based denoisers, as we show in Section 5.

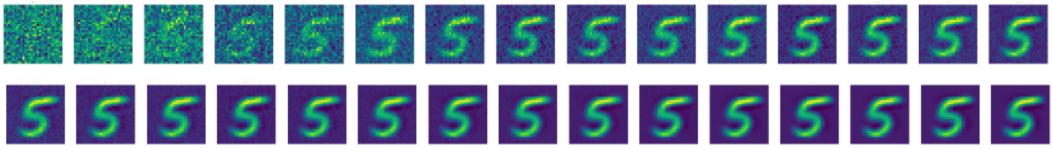

Figure 1: Digit generation from pure noise (class conditioned). The reverse process runs from left to right.

Empirical Demonstration of a Linear Diffusion Model. To illustrate the forward and backward processes in the linear case, we perform a numerical simulation using the MNIST dataset, which is simple enough to be estimated via a linear model. We start here with the training and generation procedures, and use the same setting and trained denoiser to demonstrate our findings throughout the paper.

In the following experiment we simulate the process described above using the MNIST dataset (we use the default train / test splits). In the class conditioned case, we learn a PCA denoiser with 30 components for each time step where $x_t = x_{t-1} + \epsilon_{\sigma_t}$, $\sigma_t \propto t$, and $T = 65$ iterations. Figure 1 shows a (decimated) example of digit generation from pure noise, where we apply the sequence of denoisers $D^t_{PCA} = P_t$, which will be more accurately defined in Section 4. In order to understand the reverse process, we now turn to analyze the gradual change of P_t, that might be expressed by the angle between the clean and noisy components over time.

Notations. We use A_t to denote the matrix A at time t, and a^t_i to denote the ith column of A_t.

4 Linear Diffusion as Basis Perturbation

We now turn to analyze the linear model presented above and show how the generation process can be seen as a kernel "correlation machine". Specifically, we are interested in the temporal (i.e., noise level) dependence of the denoiser Equation 8 throughout the generation process. Recall that at each time step $x_t = x_{t-1} + \epsilon_{\sigma_t}$, where $\epsilon_{\sigma_t} \sim \mathcal{N}(0, \sigma_t^2 \mathbf{I})$ (Equation 3). Since the noise is assumed to be Gaussian, we can write $x_t = x_0 + \epsilon_{\bar{\sigma}_t}$, where $\bar{\sigma}_t = \sqrt{\sum_{i=0}^{t} \sigma_i^2}$. Assume that the data distribution is such that its population covariance is given by

$$\Sigma_t = \mathbb{E} x_0 x_0^\dagger + \bar{\sigma}_t^2 \mathbf{I} = \sum_{i=0}^{r-1} \lambda_i^2 u_i u_i^\dagger + \bar{\sigma}_t^2 \mathbf{I} \triangleq \Sigma_0 + \bar{\sigma}_t^2 \mathbf{I}, \tag{9}$$

where $r - 1 < d$, i.e., the data reside in a low dimensional subspace (which is generally true for natural data). This is known as the "spiked model" (Johnstone, 2001), with a vast body of work covering the distribution and identifiability of the spikes spectrum (e.g., (Nadler, 2008)). Throughout the paper, we use the term "index" to refer to to the index i in 9, where the eigenvalues λ_i are ordered largest to smallest.

Given n samples concatenated as columns in the matrix X_0, at each time step we learn the PCA basis associated with $X_t = X_0 + E_{\bar{\sigma}_t}$, by the diagonalization of the sample covariate matrix

$$\hat{\Sigma}_t = \frac{1}{n} X_t X_t^\dagger = \frac{1}{n} (X_0 X_0^\dagger + X_0 E_{\bar{\sigma}_t}^\dagger + E_{\bar{\sigma}_t} X_0^\dagger + E_{\bar{\sigma}_t} E_{\bar{\sigma}_t}^\dagger) \triangleq U_t S_t U_t^\dagger. \tag{10}$$

Thus, during the reverse process, at each time step we apply the projection

$$D^t_{PCA} = P_t = U_t U_t^\dagger. \tag{11}$$

In order to understand the diffusion generation process, we analyze the decay of the product $\langle u^t_i, u_i \rangle$ over time, where u^t_i is the ith column of U_t. Note, that there are two drivers of change in the perturbation of u_i to u^t_i. The first being the added noise, i.e., $\|\Sigma_t - \Sigma_0\|$. This is the key in the diffusion process and our main focus. The second, is in the finite sample approximation $\left\|\hat{\Sigma}_t - \Sigma_t\right\|$. This source of error is interesting in the context of sample complexity, as it encompasses the approximation of the denoiser learned from a finite dataset, the equivalent of the sample complexity of learning the score function in (Chen et al., 2023a; Block et al., 2020; Biroli & Mézard, 2023).

For the rank-1 case, (Nadler, 2008) presented a finite sample theorem which holds with high probability for the closeness between the leading eigenvalue and eigenvector of sample and population PCA under

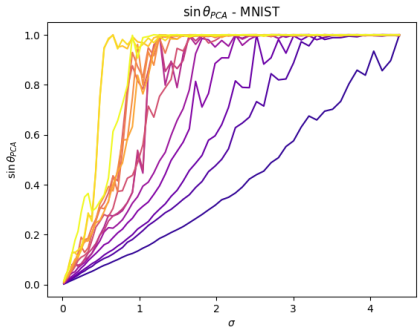

Figure 2: The sine of the angle between the clean principal components and their noisy versions, colored by the order of the eigenvalues (the darkest being largest eigenvalue). Low frequencies emerge earlier in the generation process (at higher noise levels). This motivates Assumption 4.1, that extends Equation 12 to higher ranks.

a spiked covariance model similar to Equation 9. They bound the angle between the leading empirical eigenvector and its population counterpart with approximately $\mathcal{O}(d)$ sample complexity, and a linear dependence on the noise level:

$$\mathbb{E}\sin\theta_{\text{PCA}} = \mathbb{E}\sqrt{1-\langle u^t, u\rangle^2} \approx \frac{\bar{\sigma}_t}{\lambda}\sqrt{\frac{d}{n}}, \tag{12}$$

where $\bar{\sigma}_t$ is assumed to be small and $d \gg 1$. This result shows that the leading eigenvector rotates in a rate proportional to the noise level. Our numerical experiments on the MNIST dataset (detailed in Section 4.1) show that this is a good approximation in practice, also for the rank-r case (Fig. 2).

Notice that in Equation 12 the angle is inversely linked to the eigenvalue, inferring a slower change with higher eigenvalues. In the reverse process, we gradually move from pure noise or high noise levels to smaller noise variance. Given the lower slope of the components corresponding to larger eigenvalues, we interpret the result in Fig. 2 as the earlier emergence of low frequencies in the generation process. The first component to be visible in the generated image is the one with the largest eigenvector, as it is the first one that shows a correlation in high noise levels. Throughout the generation process, when the noise level decreases, the next components take presence, by the order of their associated eigenvalue - from the larger to the smaller. Finally, the components with the smallest eigenvalues appear when the noise level is low.

In the linear case, Equation 12 shows that the diffusion model's sample complexity is determined by the sample complexity of PCA, with a linear dependence on the dimension of the data. To further enhance our understanding of the relationship between the amount of training data and the generalization of the diffusion model, we repeat the experiment with varying datasets sizes. Figure 3 shows the angle to noise profile for selected principal components, with the indices $0,5,10$ (left to right; index 0 corresponds to the largest in a list of ordered eigenvalues). Increasing the amount of training data improves robustness to noise and enables the emergence of higher frequency components at higher noise levels, thereby capturing more nuances in the generated data.

4.1 The Generated Distribution

We now turn to discuss the distribution of the generated output, and how it relates to the natural data distribution. First, we analyze a generation process by repetitive denoising without additional noise, and show how it relates to power iteration. Then we discuss a similar process only with the injection of noise, reminiscent of other common sampling methods (e..g (Ho et al., 2020)). Given our linear model, the generation process is essentially the linear transformation given by the matrix

$$\mathcal{P}_T = \Pi_{t=0}^{T} P_t = P_0 \cdots P_t \cdots P_T. \tag{13}$$

The generated output can be expressed as

$$x_g = \mathcal{P}_T \xi, \tag{14}$$

where $\xi \sim \mathcal{N}(0, \sigma_T)$. Other than the visual aesthetic of the generated images, we are interested in their distribution, and how well it represents the natural distribution of train images. Thus, we would like to compare the generated covariance $\mathbb{E}x_g x_g^{\dagger}$ to the natural covariance Σ_0.

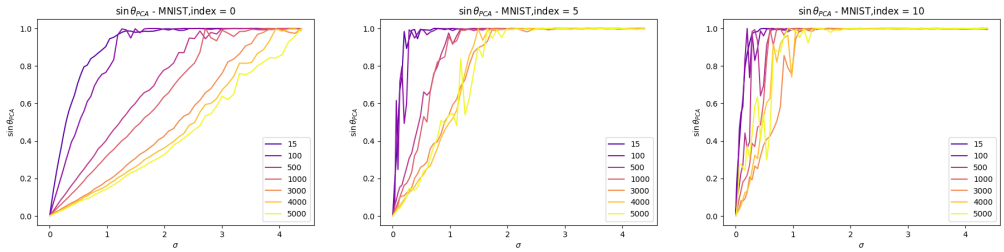

Figure 3: Effect of dataset size. The plots show $\sin\theta_{\text{PCA}}$ at different noise levels when trained on datasets with increasing size (lighter color). Each plot is of a different component index, for indices $0,5,10$ (left to right; index 0 corresponds to the largest eigenvalue). Increasing the amount of training data improves the robustness to noise, and allows the appearance of high frequencies at higher noise levels, hence capturing more data nuances in the generated data and better generalization.

In this context, a natural comparison is the power iteration (PI) method, which may be used to estimate the leading eigenvector of a matrix. This can be seen as another iterative form of generating data from random vectors. Unlike our projection, in PI we "project" a random vector onto the entire matrix, i.e. including the eigenvalues. In this case the denoiser would be $D^t_{PI} = \Sigma_0 \forall t$, where we ignore the normalization and focus on the direction of the final vector, since there is no normalization constraint for generated data in diffusion models.

We now turn to show how the reverse process performed by a repeated denoising as in Equation 13 converges in mean to PI. To this end, we make the following assumptions.

Assumption 4.1. *Assume that Equation 12 holds for all eigenvectors, i.e.,*

$$\mathbb{E}\sqrt{1-\langle u^t_i, u_i\rangle^2} \approx \frac{\bar{\sigma}_t}{\lambda_i}\sqrt{\frac{d}{n}}, \tag{15}$$

for $i=,\ldots,r-1$.

This assumption is the extension of Equation 12 to higher ranks, and is motivated by our simulations (Fig. 2). In addition, we make the following assumption regarding the cross products of components of different indices, at consecutive time steps.

Assumption 4.2. *For each index i there exists a time τ_i, where for $t \leq \tau_i$ and $j \leq i$,*

$$\mathbb{E}\langle u^t_i, u^{t+1}_j\rangle = 0. \tag{16}$$

In addition, $\tau_i > \tau_j$ for $i < j$.

This assumption is supported by our simulations in Fig. 5, and will be further discussed hereafter. Assumptions 4.2, 4.1 are an extension of Nadler (2008) to higher ranks. We leave their explicit derivation to future work, and focus on their implications to linear diffusion.

We are now ready to state our main result.

Theorem 4.3 (Convergence to Power Iteration)**.** *Let $\sigma_t = \frac{1}{T}$, $t=0,\ldots,T$. Assuming 4.2, 4.1, in the limit $T \to \infty$,*

$$\mathbb{E} x_g x_g^\dagger \propto u_0 u_0^\dagger. \tag{17}$$

Proof. Let us analyze the product in Equation 13 to show how it relates to the power method. The linear operator representing the reverse process can be written as

$$\mathcal{P}_T = U_0 \Pi_{t=0}^{T-1} (U_t^\dagger U_{t+1}) U_T^\dagger. \tag{18}$$

The matrix product $U_t^\dagger U_{t+1}$ can be analyzed using the extension of Equation 12 to higher ranks. Given 4.1, the expected inner product with the natural data component $u_i = u_i^{t=0}$ is given by

$$\mathbb{E}\langle u^t_i, u_i\rangle \approx 1 - \frac{\bar{\sigma}_t^2}{\lambda_i^2}\frac{d}{n}. \tag{19}$$

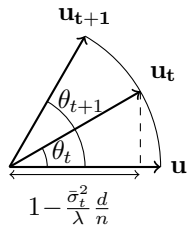

Figure 4: Schematic illustration of the basis perturbation, per index.

The evolution of this product over time is depicted in Figure 4. We are interested in the projection of u_{t+1} onto u_t, which is the cosine of the angle $\Delta\theta = \theta_{t+1} - \theta_t$. This angle is tractable for small noise levels, so we divide our analysis to two parts: $0 \le t \le \tau$ and $\tau \le t \le T$, where the choice of τ will soon be motivated.

First, we inspect the limit of $t \to 0$ ($0 \le t \le \tau$). For small angles, we can write

$$\Delta\theta = \arccos\Big(1 - \frac{\bar{\sigma}_{t+1}^2}{\lambda^2}\frac{d}{n}\Big) - \arccos\Big(1 - \frac{\bar{\sigma}_t^2}{\lambda^2}\frac{d}{n}\Big) \approx \frac{d}{\lambda^2 n}(\bar{\sigma}_{t+1}^2 - \bar{\sigma}_t^2) = \frac{\sigma_{t+1}^2 d}{\lambda^2 n}, \tag{20}$$

since $\arccos\theta \approx \frac{\pi}{2} - \theta$ and $\bar{\sigma}_t^2 = \sum_{\tau=0}^{t} \sigma_\tau^2$. The diagonal elements in $U_t^\dagger U_{t+1}$ are then given by

$$\mathbb{E}\langle u_i^t, u_i^{t+1}\rangle \approx \cos\frac{\sigma_{t+1}^2 d}{\lambda_i^2 n}, \tag{21}$$

where the off-diagonal elements are negligible, since

$$\mathbb{E}\langle u_i^t, u_j^{t+1}\rangle \approx \mathbb{E}\langle u_i^t, u_j^t\rangle = 0, \tag{22}$$

which holds for $t \le \tau_{r-1}$ by Assumption 4.2. Notice, that in small angles, $\langle u_i^t - u_i^{t+1}, u\rangle = (\sigma_{t+1}^2 d)/(\lambda^2 n) \to 0$, so the vectors u_i^t are co planar, as depicted in Figure 4. Thus, the time point basis correlations $U_t^\dagger U_{t+1}$ form an approximately diagonal matrix with the fraction $c_i \triangleq \cos\frac{\sigma_{t+1}^2 d}{\lambda_i^2 n}$ on the diagonal, where $c_i > c_j$ for $i < j$. We eliminate the dependence of c_i on t by choosing the constant schedule $\sigma_t = 1/T\ \forall t$, to simplify the proof. However, many schedules can be used, as long as $c_{i,t} > c_{j,t}$ remains correct. Define the partial linear diffusion operator until time τ by $\mathbb{E}\mathcal{P}_\tau = \Pi_{t=0}^{\tau} P_t$. Then

$$\mathbb{E}\mathcal{P}_\tau = U_0 \begin{pmatrix} c_0^\tau & & \\ & \ddots & \\ & & c_{r-1}^\tau \end{pmatrix} U_\tau^\dagger = U_0 c_0^\tau \begin{pmatrix} 1 & & \\ & \left(\frac{c_1}{c_0}\right)^\tau & \\ & & \ddots \end{pmatrix} U_\tau^\dagger \underset{\tau\to T}{\to} U_0 \begin{pmatrix} c_0^\tau & & \\ & 0 & \\ & & \ddots \end{pmatrix} U_\tau^\dagger, \tag{23}$$

where the diagonal elements decay as τ grows larger, as $c_i > c_j$ for $i < j$. Similarly to power iteration, the convergence rate depends on the ratio c_1/c_0. The convergence rate might not be fast enough for the process to converge while the small angles approximation still holds. Thus, we continue with the second phase of our analysis, showing the convergence of the full reverse process.

We now turn to analyse the phase where $\tau \le t \le T$. In high noise levels, the correlation with the natural basis is low, and the products $U_t^\dagger U_{t+1}$ are not exactly diagonal. However, the correlation "leaks" to a close neighborhood of the original component and the temporal products are still somewhat concentrated around their diagonal. This process happens in accordance with Equation 4.1, where the spreading of the diagonal elements happens for high indices in lower values of t (less noise is needed to spread the correlation). This leads us to Assumption 4.2, claiming that for each index i there exists a time τ_i after which the small angle approximation does not hold; $\tau_i > \tau_j$ for $i < j$. This is apparent in practice, and depicted in 5 (left image per duo). However, given the decaying diagonal structure of the partial operator $\mathcal{P}_\tau$, we will now show that 4.2 is sufficient for the total operator to converge as desired.

Suppose we added one more matrix multiplication to our former analysis, i.e. observe

$$\mathbb{E}\mathcal{P}_\tau U_{\tau+1} = U_0 c_0^\tau \begin{pmatrix} 1 & & \\ & \left(\frac{c_1}{c_0}\right)^\tau & \\ & & \ddots \end{pmatrix} U_\tau^\dagger U_{\tau+1}. \tag{24}$$

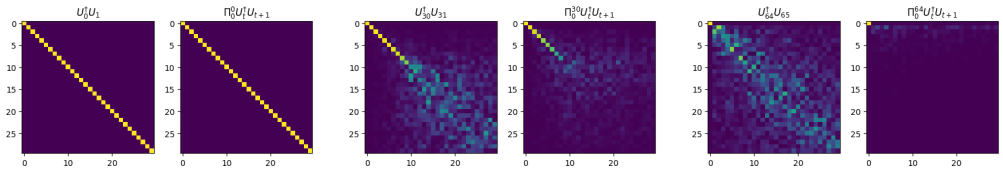

Figure 5: The time point basis correlation matrices $U_\tau^T U_{\tau+1}$ (left per pair), together with the partial product $\Pi_{t=0}^{\tau}(U_t^\dagger U_{t+1})$ (right per pair) at different time points. This justifies Assumption 4.2, and shows that the total projection (bottom right image, for $\tau = T$) converges to the first eigenvector, similarly to the power method.

Assumption 4.2 guarantees $U_\tau^\dagger U_{\tau+1}$ is diagonal just enough not to spoil the diagonality of the next partial operator $\mathbb{E}\mathcal{P}_{\tau+1}$. To see this, let us inspect some intermediate index i, where the entries in $j > i$ are already practically zero. Thus, we have

$$\mathbb{E}\mathcal{P}_{\tau_i} U_{\tau_i+1} = U_0 c_0^{\tau_i+1} \underbrace{\begin{pmatrix} 1 & & & \\ & \ddots & & \\ & & \left(\frac{c_i}{c_0}\right)^{\tau_i} & \\ & & & \mathbb{O} \end{pmatrix}}_{\triangleq C_{\tau_i}} \begin{pmatrix} 1 & & & \\ & \ddots & & \\ & & \frac{c_i}{c_0} & \\ & & & \mathbb{A} \end{pmatrix} = U_0 c_0^{\tau_i+1} \begin{pmatrix} 1 & & & \\ & \ddots & & \\ & & \left(\frac{c_i}{c_0}\right)^{\tau_i+1} & \\ & & & \mathbb{O} \end{pmatrix}$$

where $\mathbb{O}$ is a block of zeros and $\mathbb{A}$ is a block matrix the same size as $\mathbb{O}$, that can have nonzero entries, by Assumption 4.2. Since the elements of the partial product C_{τ_i} decay faster with i than any single product $U_{\tau_i}^\dagger U_{\tau_i+1}$, C_{τ_i+1} is also diagonal. Overall, the final product is a diagonal matrix with a spectrum that converges to be concentrated around the first eigenvalue, where we can control the distribution of the generated data by the choice of the diffusion parameters. Figure 5 shows our simulation of the process, supporting both assumption 4.2 and the result stated by this theorem. □

Oftentimes, the reverse process includes the injection of noise to the intermediate images (e.g. (Ho et al., 2020)). The overall transformation in this case is given by

$$x_g = \Sigma_{t=0}^{T} \Pi_{\tau=0}^{t} P_\tau \xi_t = P_0 \cdots P_T \xi_T + \cdots + P_0 \xi_0 \tag{25}$$

for some schedule $\{\xi_t\}_{t=0}^{T}$ (for example, $\xi_T \sim \mathcal{N}(0,1)$ and $\xi_t = \mathcal{N}(0,1/T)$ for $t = 0,\ldots,T-1$). In this case, the generated output is a combination of a (purely) noisy image that was repeatedly correlated to converge to v_0 (as shown above), with generally lower noise levels that are "lightly" correlated, although to the cleaner projection operators. The generated output can thus be seen as a combination of three conceptual parts, with a different balance of the noise level and the portrayed components.

The first eigenvector The first part of the sum in Equation 25 is $P_0 \cdots P_T \xi_T$, the estimation of the eigenvector with the largest eigenvalue, as shown above theoretically in Equation 23 and empirically in the right matrix of the bottom right duo in Fig. 5. The "strongest" noise is repeatedly correlated to be concentrated around the first eigenvector.

The entire (clean) spectrum The last part in Equation 25 is $P_0 \xi_0$, a weak noise level that is spread across all components. This noise is very lightly and not repeatedly correlated, although to a clean version of the natural data basis.

In between The third part consists of all the intermediate products $\Pi_{\tau=0}^{t} P_\tau \xi_t$. The product operators $\Pi_{\tau=0}^{t} P_\tau$ preserve varying parts of the natural spectrum, according to t - as t grows, the total projection tends to retain only the components associated with larger eigenvalues. This can bee seen in Fig. 5. The right matrix in each pair shows the product $\Pi_{\tau=0}^{t} P_\tau$ for varying values of t. The total projections range from the entire spectrum (top left), to only the leading eigenvalue (bottom right). In between, the products are diagonal matrices where the entries in the indices of the smaller eigenvalues have already diminished, in a similar way to the convergence described in Equation 23.

Thus, we get a combination of a solid estimation of the leading eigenvector, together with a more uniform and week sampling of the components with low eigenvalues in the natural data basis. In between, the intermediate projections are at different levels of convergence to the leading eigenvector, hence tend to be more concentrated on components with large eigenvalues as $t \to T$. The freedom in choice of schedule $\{\xi_t\}_{t=0}^{T}$, allows control of the spread of the final distribution on the natural data components.

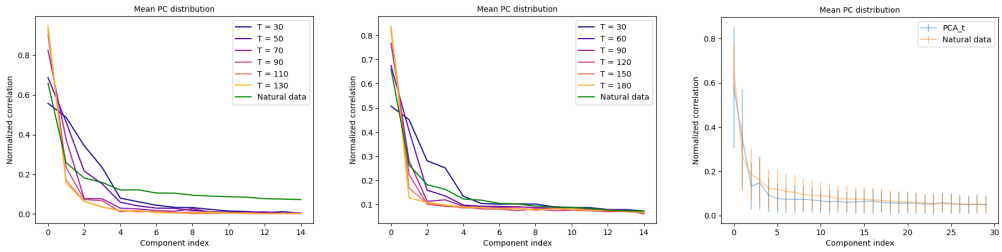

Figure 6: The empirical distribution of generated images over the natural principal components, with (middle) and without (left) injected noise. On the right - the best configuration with the generated standard deviation (see Sec. 4.1).

To inspect this, we plot the empirical distribution of generated images over the clean PCs, given by

$$p_i = \frac{1}{n}\sum_{j=1}^{n}\frac{|\langle u_i, x_j\rangle|}{\|x_j\|_2}, \tag{26}$$

where u_i is the clean principal component with index i (defined in Equation 9) and x_j is a generated sample, out of n examples. Figure 6 shows the empirical distribution of generated images over the clean principal components. On the left, we plot the distribution without injected noise, i.e. $x_g = \mathcal{P}_T \xi$ (as in Equation 14), for various values of T. As we show above, the distribution tends to be concentrated on the first eigenvector as T increases. The center plot shows the distribution of the process including the injected noise in the intermediate denoising steps. While in the low indices the dominant behavior is similar to the former case, the higher orders do not converge to zero and maintain their presence in the generated distribution. We note, that more sophisticated nonlinear deterministic samplers might not require the injection of noise in order to converge to the natural data distribution (e.g. Lu et al. (2022)). However, given a linear model, it is natural to accept added stochasticity in the lack of nonlinearity (more on that in Section 5). On the right, we picked the best configuration ($T = 65$ in this case) to approximate the natural distribution. Notice, that the final generated distribution depends on the choice of parameters, where one can control the mean of the generated spectrum (this might be a feature for some applications, such as segmentation via diffusion, etc.). It might be interesting to derive the optimal parametrization for the convergence of the linear model - we leave this for future work. In addition to the convergence in mean, we included the standard deviation of the natural and generated samples, resulting in a decent fit to the target distribution.

5 Empirical Extension to Deep Denoisers

In the linear case described above, the optimal denoiser is given by the PCA projection onto the clean(er) data. These denoisers are computed with the training data, and their principal components do not depend on the input in the reverse process. When the denoiser is nonlinear, and might be implemented using a deep neural network, its input-output mapping can be locally expressed via the network Jacobian, by

$$D(x_t) = \nabla D(x_t) x_t = V_t \Lambda_t V_t^{\dagger} x_t, \tag{27}$$

where $V_t \Lambda_t V_t^{\dagger}$ denotes the eigen decomposition of the Jacobian calculated at x_t. For simplicity, we assume that the Jacobian is symmetric and non-negative (which is approximately true (Mohan et al., 2020)). Note that in this case, the denoising base depends on the input image (and noise level). While the network is non linear, we can follow the generation path in the sampling process and inspect the basis of the network Jacobians calculated at the intermediate sampled points x_t. We can then trace $\sin\theta_J = \sqrt{1 - \langle v_i^t, v_i^{t=0}\rangle^2}$ where the subscript "J" stands for Jacobian, v_i^t is the ith column in V_t defined in Equation 27, in a similar way to our simulations of the linear case (Figure 2). This can be calculated per generation path, where x_0 is the final generated image, and V_0 is the basis of the Jacobian calculated at this final point.

Figure 7 shows $\sin\theta_J$ calculated using the Jacobians of a UNet based diffusion model, described in (Ning et al., 2023). This model was simply chosen as the [1]state-of-the-art in the task of image generation considering the CelebA dataset at the time of writing this paper. We used the default settings and calculated the Jacobians at the final iterations. We plot the results for the leading 300 Jacobian eigenvectors, where the color is assigned by the index - darker colors for lower indices i. We repeated the experiment sampling images from the CelebA dataset (left) and CIFAR 10 (right). Even though the denoising model is far from linear, the decay of the angle between the denoising basis in high noise levels and the natural denoising basis is similar to the decay in the linear case (compare to Figure 2). In this case as well, the correlation of the low indices (and hence low frequencies) withstands higher noise levels, thus appearing first in the generation process. As this is the basis of our analysis comparing the reverse diffusion process to power iteration, this experiment shows that our analysis is relevant in a broader context and not just in the simplified linear case.

This analysis focuses on the local behavior of the nonlinear denoiser at the end of the generation process, demonstrating its similarity to a linear denoising chain. Each plot represents a single generation path, not the overall distribution of generated outputs.

While linear diffusion models are easy to analyze, they may struggle to generate complex datasets. Nonlinear models, on the other hand, can navigate a diverse set of linearized regions during the generation process (as illustrated in Figure 7). This allows them to generate diverse data even without added noise, unlike linear models which ultimately converge in mean to a single point (Theorem 4.3) and therefore require noise injection for diverse outputs. This contrasts with some deterministic nonlinear samplers (e.g., Lu et al. (2022)) that do not rely on added noise.

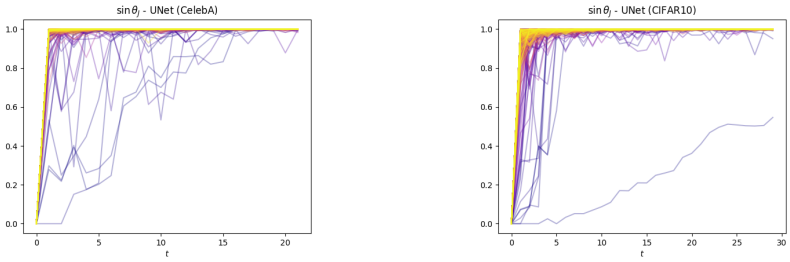

Figure 7: Image generation - the sine of the angle between Jacobian eigenvectors at the final generated image ($t=0$) and intermediate iterations ($t>0$). The diffusion model includes a UNet-based denoiser trained on CelebA (left) or CIFAR10 (rignt). Color by index (the darker the color the lower the index, referring to columns of the Jacobian basis V_t). The Jacobians of the nonlinear denoiser conform to the behavior of the linear model.

6 Conclusion

In this paper, we discuss a simple diffusion model with a linear denoiser and normalization free sampler, that allows us to cast the diffusion problem as noisy PCA, and make the connection to the spiked covariance model assuming that the natural data distribution reside in a low dimensional subspace. This enables us to show that in the linear case, the generation process acts as a "correlation machine", where initial random noise is repeatedly correlated to noisy estimations of the natural data basis, to finally embody the true distribution, in a manner similar to the power iteration method. We show that in this process, low frequencies emerge earlier, and more data contributes to a richer representation per the same diffusion configuration. Finally, we demonstrate the relevance of our analysis also in a deep, non-linear diffusion denoiser.

We acknowledge the limitation of admitting a linear model, with its lack of ability to represent the complex data often expected of diffusion models. While our theoretical setting is modest, we empirically demonstrate how our observations deduced from a simple linear model and classic theory (Johnstone, 2001; Nadler, 2008) are relevant to more general models and datasets. This enables us to shed light on the internal mechanism powering this technology, and connect it to a rich pool of theory and prevalent methods such as power iteration.

[1]https://paperswithcode.com/sota/image-generation-on-celeba-64x64

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A POWER ITERATION AND ITS CONVERGENCE

Power Iteration is a simple algorithm used to compute the dominant eigenvalue and its corresponding eigenvector of a matrix. It iteratively refines an initial random vector by multiplying it with the matrix, which gradually aligns with the eigenvector corresponding to the largest eigenvalue. For a thorough introduction to the method, see e.g. Andrilli & Hecker (2023). Given a square matrix $A \in \mathbb{R}^{n \times n}$, the goal is to compute the dominant eigenvalue λ_1 and its corresponding eigenvector v_1. The Power Iteration algorithm is defined as follows:

Algorithm 1 Power Iteration Algorithm

Input: Matrix $A \in \mathbb{R}^{n \times n}$, initial vector $v_0 \in \mathbb{R}^n$, number of iterations k
Output: Approximate dominant eigenvector v_k
Normalize the initial vector: $v_0 \leftarrow \frac{v_0}{\|v_0\|}$
for each iteration $i = 1,2,...,k$ **do**
  $v_i \leftarrow A v_{i-1}$
  Normalize $v_i \leftarrow \frac{v_i}{\|v_i\|}$
end for
return v_k

The algorithm starts with an arbitrary vector v_0, which is normalized to ensure numerical stability. In each iteration, the vector v_i is updated by multiplying it with the matrix A, followed by normalization. After k iterations, the vector v_k is expected to be close to the eigenvector corresponding to the largest eigenvalue of A.

A.1 CONVERGENCE ANALYSIS

Let A be a square matrix with eigenvalues $\lambda_1, \lambda_2, ..., \lambda_n$, where the eigenvalues are ordered such that $|\lambda_1| > |\lambda_2| \geq \cdots \geq |\lambda_n|$. Denote the corresponding eigenvectors by $v_1, v_2, ..., v_n$, where v_1 is the eigenvector corresponding to the dominant eigenvalue λ_1.

The key idea behind Power Iteration is that, after sufficient iterations, the sequence of vectors v_i converges to the eigenvector associated with λ_1, under certain conditions.

Let v_0 be the initial vector, which can be expressed as a linear combination of the eigenvectors of A:

$$v_0 = \sum_{i=1}^{n} \alpha_i v_i$$

where α_i are scalar coefficients. After applying the matrix A in each iteration, we obtain the sequence of vectors:

$$v_i = A v_{i-1} = A \left(\sum_{i=1}^{n} \alpha_i v_i \right) = \sum_{i=1}^{n} \alpha_i \lambda_i^i v_i$$

Thus, the i-th iteration amplifies the component of v_0 along the direction of the eigenvector corresponding to the eigenvalue λ_1, while the other components decay at a rate proportional to the magnitude of their respective eigenvalues. As the iterations proceed, the contribution of the eigenvectors associated with smaller eigenvalues diminishes, and the vector v_i becomes increasingly aligned with v_1, the eigenvector corresponding to λ_1.

Formally, we express the evolution of v_i as:

$$v_i = \lambda_1^i \alpha_1 v_1 + \lambda_2^i \alpha_2 v_2 + \cdots + \lambda_n^i \alpha_n v_n$$

The relative influence of the eigenvectors corresponding to $\lambda_2, \lambda_3, ..., \lambda_n$ decays exponentially as $i \to \infty$ because $\lambda_1 > |\lambda_2| \geq \cdots \geq |\lambda_n|$. Specifically, the error in approximating v_1 decreases at a rate proportional to $\frac{|\lambda_2|}{|\lambda_1|}$, leading to the following convergence result:

$$\frac{\|v_i - \lambda_1^i v_1\|}{\|v_1\|} \leq C \left(\frac{|\lambda_2|}{|\lambda_1|} \right)^i$$

for some constant C, where $\|\cdot\|$ is the vector norm (usually the Euclidean norm).

Therefore, the Power Iteration algorithm converges to the dominant eigenvector v_1 at a rate determined by the ratio of the magnitudes of the first and second largest eigenvalues, $\rho = \frac{|\lambda_2|}{|\lambda_1|}$. If λ_2 is much smaller than λ_1, convergence is fast. However, if λ_2 is close to λ_1, convergence can be slow, requiring more iterations to achieve a satisfactory approximation. The convergence is linear, with the error decaying exponentially as the number of iterations increases. For a matrix A with a well-separated dominant eigenvalue λ_1 (i.e., $|\lambda_1| \gg |\lambda_2|$), Power Iteration converges quickly, typically in $O(\log(\epsilon)/\log(\rho))$ iterations to achieve an error of size ϵ.