

DEPARTMENT OF COMPUTER SCIENCE

Video Diffusion Models for Climate Simulations

	George Herbert	
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Abstract

Dedication and Acknowledgements

Declaration

I declare that the work in this dissertation was carried out in accordance with the requirements of the University's Regulations and Code of Practice for Taught Programmes and that it has not been submitted for any other academic award. Except where indicated by specific reference in the text, this work is my own work. Work done in collaboration with, or with the assistance of others, is indicated as such. I have identified all material in this dissertation which is not my own work through appropriate referencing and acknowledgement. Where I have quoted or otherwise incorporated material which is the work of others, I have included the source in the references. Any views expressed in the dissertation, other than referenced material, are those of the author.

George Herbert, Wednesday $19^{\rm th}$ April, 2023

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Ethics Statement

Notation and Acronyms

i.i.d. : Independent and identically distributed

 KL : Kullback–Leibler

VAE : Variational Autoencoder

:

 $\mathbf{X}^{\circ n}$: Element-wise exponentiation of matrix \mathbf{X} with power n diag(\mathbf{x}) : Diagonal matrix with the values of vector \mathbf{x} on the diagonal

 $\log(x)$: Natural logarithm function (i.e. logarithm with base e) applied to x

 $f \simeq g$: g is an unbiased estimator of f

Chapter 1

Introduction

Chapter 2

Technical Background

2.1 Generative Models

Let us consider some dataset \mathcal{D} consisting of $N_{\mathcal{D}} \geq 1$ datapoints which we assume are independent and identically distributed (i.i.d.):

$$\mathcal{D} = \{ \mathbf{x}^{(i)} \mid 1 \le i \le N_{\mathcal{D}}, i \in \mathbb{N} \}$$

$$(2.1)$$

We assume each observed datapoint $\mathbf{x}^{(i)} \in \mathcal{D}$ is a realisation of the observed random variable \mathbf{x} from an underlying process, whose true distribution $p^*(\mathbf{x})$ is unknown. We will omit the index (i) whenever it is clear we are referring to a single datapoint. The goal of *generative modelling* is to approximate this true distribution with a chosen model $p_{\theta}(\mathbf{x})$ with parameters θ . We learn parameters θ such that the probability distribution function given by the model $p_{\theta}(\mathbf{x})$ approximates the true distribution of the data, such that for any observed \mathbf{x} :

$$p_{\theta}(\mathbf{x}) \approx p^*(\mathbf{x})$$
 (2.2)

Once learnt, we can generate new samples *unconditionally* from our approximate model at will. We thus refer to the model $p_{\theta}(\mathbf{x})$ as an unconditional generative model.

2.2 Conditional Generation

We can extend generative modelling to the conditional setting. We consider each observed \mathbf{x} to have some corresponding conditioning information \mathbf{c} . In this context, we wish to approximate the conditional distribution $p^*(\mathbf{x}|\mathbf{c})$. Similar to the unconditional setting, we learn parameters θ for our model $p_{\theta}(\mathbf{x}|\mathbf{c})$ such that for any observed \mathbf{x} and conditioning information \mathbf{c} :

$$p_{\theta}(\mathbf{x}|\mathbf{c}) \approx p^*(\mathbf{x}|\mathbf{c})$$
 (2.3)

Once learnt, we can generate new samples *conditionally* from our approximate model at will.

One of the most basic cases is a class-conditional generative model, where the conditioning variable \mathbf{c} is simply a class label. In such cases, our conditional model $p_{\theta}(\mathbf{x}|\mathbf{c})$ has an interpretation as the reverse of a discriminative classification model—a more traditional form of machine learning. As opposed to inputting an observed \mathbf{x} and the model outputting the predicted corresponding class label \mathbf{c} , we input a class label \mathbf{c} and use the model to generate a new sample $\mathbf{x} \sim p_{\theta}(\mathbf{x}|\mathbf{c})$.

Significantly, the conditioning variable \mathbf{c} is not limited to class labels; it can be flexible and take the form of any additional information we wish to condition on to generate samples. A powerful tool in the case of image generation, \mathbf{c} may be a text encoding to facilitate text-to-image synthesis (see e.g. [12, 6]). Alternatively, \mathbf{c} may be a lower-resolution image from which we wish to upscale to add higher-resolution details, known as image super-resolution (see e.g. [4]).

In this work, we do not explicitly use a conditional model. However, we do derive one approximately from an unconditional model. We discuss this further in Section 2.7.10.

2.3 Latent Variables

We can think of each observed datapoint $\mathbf{x} \in \mathcal{D}$ as being represented or generated via $N_{\mathbf{z}} \geq 1$ associated latent variables:

$$\{\mathbf{z}_1, \dots, \mathbf{z}_{N_{\mathbf{z}}}\} = \{\mathbf{z}_i \mid i < 1 < N_{\mathbf{z}}, i \in \mathbb{N}\}$$

$$(2.4)$$

The latent variables are part of the model, but we do not observe them directly, and they are not within the dataset. We model the joint distribution of the observed variable and the latent variables by $p_{\theta}(\mathbf{x}, \mathbf{z}_1, \dots, \mathbf{z}_{N_z})$; the marginal distribution over the observed variable $p_{\theta}(\mathbf{x})$ is given by:

$$p_{\theta}(\mathbf{x}) = \int p_{\theta}(\mathbf{x}, \mathbf{z}_1, \dots, \mathbf{z}_{N_{\mathbf{z}}}) d\mathbf{z}$$
 (2.5)

In the context of a latent-variable model, generation refers to the process of sampling the latent variables from the joint distribution $p_{\theta}(\mathbf{z}_1, \dots, \mathbf{z}_{N_z})$, and then sampling the observed variable from the conditional distribution $p_{\theta}(\mathbf{x}|\mathbf{z}_1, \dots, \mathbf{z}_{N_z})$. In the simplest case, we may only have a single latent variable; we omit the index 1 in such cases for notational simplicity.

2.4 Likelihood-Based Generative Models

As mentioned in Section 2.1, the goal of a generative model is to learn parameters θ such that $p_{\theta}(\mathbf{x}) \approx p^*(\mathbf{x})$. One way to interpret this is as a minimisation problem. Namely, we wish to learn parameters θ that minimise the Kullback–Leibler (KL) divergence of the true distribution $p^*(\mathbf{x})$ from our model distribution $p_{\theta}(\mathbf{x})$:

$$\underset{\theta}{\operatorname{arg\,min}} \ D_{KL}(p^*(\mathbf{x}) || p_{\theta}(\mathbf{x})) \tag{2.6}$$

The KL divergence, denoted D_{KL} , measures the dissimilarity between two probability distributions; in our case, it provides a measure of the information lost when we approximate the true distribution $p^*(\mathbf{x})$ with our model distribution $p_{\theta}(\mathbf{x})$.

We can reformulate the KL divergence of the true distribution $p^*(\mathbf{x})$ from our model distribution $p_{\theta}(\mathbf{x})$ to provide a likelihood-based interpretation:

$$D_{KL}(p^*(\mathbf{x})||p_{\theta}(\mathbf{x})) = \mathbb{E}_{\mathbf{x} \sim p^*(\mathbf{x})} \left[\log \left(\frac{p^*(\mathbf{x})}{p_{\theta}(\mathbf{x})} \right) \right]$$
(2.7)

$$= \mathbb{E}_{\mathbf{x} \sim p^*(\mathbf{x})} \left[\log p^*(\mathbf{x}) \right] + \mathbb{E}_{\mathbf{x} \sim p^*(\mathbf{x})} \left[-\log p_{\theta}(\mathbf{x}) \right]$$
 (2.8)

$$= -\mathcal{H}(p^*(\mathbf{x})) + \mathbb{E}_{\mathbf{x} \sim p^*(\mathbf{x})} \left[-\log p_{\theta}(\mathbf{x}) \right]$$
 (2.9)

where $\mathcal{H}(p^*(\mathbf{x}))$ is the entropy of the true distribution $p^*(\mathbf{x})$ and is constant. As such, minimisation of the KL divergence in this context equates to minimisation of the expected negative log-likelihood of our model distribution $p_{\theta}(\mathbf{x})$ with respect to $\mathbf{x} \sim p^*(\mathbf{x})$; formally:

$$\underset{\theta}{\operatorname{arg\,min}} \ D_{KL}(p^*(\mathbf{x}) \| p_{\theta}(\mathbf{x})) = \underset{\theta}{\operatorname{arg\,min}} \ \mathbb{E}_{\mathbf{x} \sim p^*(\mathbf{x})} \left[-\log p_{\theta}(\mathbf{x}) \right]$$
 (2.10)

Under the assumption that each of the $N_{\mathcal{D}}$ samples in our dataset \mathcal{D} are i.i.d. according to $p^*(\mathbf{x})$, we can construct an unbiased estimator:

$$\mathbb{E}_{\mathbf{x} \sim p^*(\mathbf{x})} \left[-\log p_{\theta}(\mathbf{x}) \right] \simeq \frac{1}{N_{\mathcal{D}}} \left(-\log p_{\theta}(\mathcal{D}) \right) = \frac{1}{N_{\mathcal{D}}} \sum_{\mathbf{x} \in \mathcal{D}} \left(-\log p_{\theta}(\mathbf{x}) \right)$$
(2.11)

In other words, under the i.i.d assumption of \mathcal{D} , the mean negative log-likelihood of our model with respect to \mathcal{D} serves as an unbiased estimator of the expected negative log-likelihood of our model with respect to $\mathbf{x} \sim p^*(\mathbf{x})$. In practice, for computational efficiency reasons—as well as GPU memory limitations—we learn via mini-batches $\mathcal{M} \subset \mathcal{D}$ of size $N_{\mathcal{M}} < N_{\mathcal{D}}$, which is itself an unbiased estimator:

$$\frac{1}{N_{\mathcal{D}}} \left(-\log p_{\theta}(\mathcal{D}) \right) \simeq \frac{1}{N_{\mathcal{M}}} \left(-\log p_{\theta}(\mathcal{M}) \right) = \frac{1}{N_{\mathcal{M}}} \sum_{\mathbf{x} \in \mathcal{M}} \left(-\log p_{\theta}(\mathbf{x}) \right) \tag{2.12}$$

As such, by transitivity, the mean negative log-likelihood of our model with respect to each mini-batch \mathcal{M} is itself an unbiased estimator of the expected negative log-likelihood of our model with respect to $\mathbf{x} \sim p^*(\mathbf{x})$. We refer to the broad class of generative models trained to minimise the expected negative log-likelihood of $p_{\theta}(\mathbf{x})$ with respect to $\mathbf{x} \sim p^*(\mathbf{x})$ as likelihood-based generative models.

2.5 Variational Autoencoders

2.5.1 Components of the Basic Variational Autoencoder

The variational autoencoder (VAE) [8, 11] is an important example of a likelihood-based generative model. In its simplest form, the VAE is a latent-variable model $p_{\theta}(\mathbf{x}, \mathbf{z})$ with a single latent \mathbf{z} . We assume that each observed datapoint \mathbf{x} is generated via a two-step process. First, a latent \mathbf{z} is generated from some true prior distribution $p^*(\mathbf{z})$, followed by an observed value \mathbf{x} generated from some true conditional distribution $p^*(\mathbf{x}|\mathbf{z})$. Thus, our model $p_{\theta}(\mathbf{x}, \mathbf{z})$ we seek to optimise such that $p_{\theta}(\mathbf{x}) \approx p^*(\mathbf{x})$ takes the following factorised form:

$$p_{\theta}(\mathbf{x}, \mathbf{z}) = p_{\theta}(\mathbf{z})p_{\theta}(\mathbf{x}|\mathbf{z}) \tag{2.13}$$

where, naturally, we need to specify our two distributions: $p_{\theta}(\mathbf{z})$ and $p_{\theta}(\mathbf{x}|\mathbf{z})$. We refer to $p_{\theta}(\mathbf{z})$ as the prior over \mathbf{z} , and one common choice is the standard Gaussian:

$$p_{\theta}(\mathbf{z}) = \mathcal{N}(\mathbf{z}; \mathbf{0}, \mathbf{I}) \tag{2.14}$$

Furthermore, we refer to $p_{\theta}(\mathbf{x}|\mathbf{z})$ as the stochastic *decoder* since given a latent \mathbf{z} it produces a distribution over the possible corresponding values of \mathbf{x} . As an example, we can select $p_{\theta}(\mathbf{x}|\mathbf{z})$ be a multivariate Gaussian with diagonal covariance:

$$p_{\theta}(\mathbf{x}|\mathbf{z}) = \mathcal{N}(\mathbf{x}; \boldsymbol{\mu}_{\theta}(\mathbf{z}), \operatorname{diag}(\boldsymbol{\sigma}_{\theta}(\mathbf{z}))^{\circ 2})$$
 (2.15)

where $\mu_{\theta}(\mathbf{z})$ and $\sigma_{\theta}(\mathbf{z})$ are outputs from a neural network with parameters θ . One final, crucial defining feature of the VAE is the stochastic encoder $q_{\phi}(\mathbf{z}|\mathbf{x})$, also referred to as the inference model, with variational parameters ϕ . The stochastic encoder $q_{\phi}(\mathbf{z}|\mathbf{x})$ approximates the intractable posterior $p_{\theta}(\mathbf{z}|\mathbf{x})$ of the generative model. Again, a common choice is for $q_{\phi}(\mathbf{z}|\mathbf{x})$ to be a multivariate Gaussian with diagonal covariance:

$$q_{\phi}(\mathbf{z}|\mathbf{x}) = \mathcal{N}(\mathbf{z}, \boldsymbol{\mu}_{\phi}(\mathbf{z}), \operatorname{diag}(\boldsymbol{\sigma}_{\phi}(\mathbf{z}))^{\circ 2})$$
 (2.16)

Figure 2.1 provides a graphical depiction of the VAE.

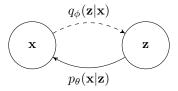


Figure 2.1: Graphical depiction of basic VAE with one observed variable \mathbf{x} and one latent variable \mathbf{z} . Solid lines depict the Bayesian network of the generative model; dashed lines depict the Bayesian network of the approximate inference model.

2.5.2 Evidence Lower Bound Objective

The VAE falls into the broad class of likelihood-based generative models. However, the likelihood $p_{\theta}(\mathbf{x})$ cannot be optimised directly, as the VAE model does not make common simplifying assumptions about marginal and posterior probabilities. Notably, we assume the model's marginal likelihood $p(\mathbf{x})$ given by:

$$p_{\theta}(\mathbf{x}) = \int p_{\theta}(\mathbf{x}|\mathbf{z})p_{\theta}(\mathbf{z})d\mathbf{z}$$
 (2.17)

does not have an analytic solution or efficient estimator. In addition, we assume the model's posterior density $p_{\theta}(\mathbf{z}|\mathbf{x})$ is intractable, so we cannot employ the expectation-maximisation algorithm.

Not making these simplifying assumptions is the reason for introducing the inference model $q_{\phi}(\mathbf{z}|\mathbf{x})$ to approximate the model's intractable posterior $p_{\theta}(\mathbf{z}|\mathbf{x})$. With the introduction of the inference model,

we can derive a variational bound on the negative log-likelihood:

$$-\log p_{\theta}(\mathbf{x}) = \mathbb{E}_{\mathbf{z} \sim q_{\phi}(\mathbf{z}|\mathbf{x})} \left[-\log p_{\theta}(\mathbf{x}) \right]$$
(2.18)

$$= \mathbb{E}_{\mathbf{z} \sim q_{\phi}(\mathbf{z}|\mathbf{x})} \left[-\log \left(\frac{p_{\theta}(\mathbf{x}, \mathbf{z})}{p_{\theta}(\mathbf{z}|\mathbf{x})} \right) \right]$$
 (2.19)

$$= \mathbb{E}_{\mathbf{z} \sim q_{\phi}(\mathbf{z}|\mathbf{x})} \left[-\log \left(\frac{p_{\theta}(\mathbf{x}, \mathbf{z}) q_{\phi}(\mathbf{z}|\mathbf{x})}{p_{\theta}(\mathbf{z}|\mathbf{x}) q_{\phi}(\mathbf{z}|\mathbf{x})} \right) \right]$$
(2.20)

$$= \mathbb{E}_{\mathbf{z} \sim q_{\phi}(\mathbf{z}|\mathbf{x})} \left[-\log \left(\frac{p_{\theta}(\mathbf{x}, \mathbf{z})}{q(\mathbf{z}|\mathbf{x})} \right) \right] - \mathbb{E}_{\mathbf{z} \sim q_{\phi}(\mathbf{z}|\mathbf{x})} \left[\log \left(\frac{q(\mathbf{z}|\mathbf{x})}{p_{\theta}(\mathbf{z}|\mathbf{x})} \right) \right]$$
(2.21)

$$= \mathbb{E}_{\mathbf{z} \sim q_{\phi}(\mathbf{z}|\mathbf{x})} \left[-\log \left(\frac{p_{\theta}(\mathbf{x}, \mathbf{z})}{q(\mathbf{z}|\mathbf{x})} \right) \right] - D_{KL}(q_{\phi}(\mathbf{z}|\mathbf{x}) || p_{\theta}(\mathbf{z}|\mathbf{x}))$$
(2.22)

The second term in Equation 2.22 is the KL divergence of $q_{\phi}(\mathbf{z}|\mathbf{x})$ and $p_{\theta}(\mathbf{z}|\mathbf{x})$ and is non-negative:

$$D_{KL}(q_{\phi}(\mathbf{z}|\mathbf{x})||p_{\theta}(\mathbf{z}|\mathbf{x})) \ge 0 \tag{2.23}$$

and zero if and only if $q_{\phi}(\mathbf{z}|\mathbf{x}) = p_{\theta}(\mathbf{z}|\mathbf{x})$. The first term in Equation 2.22 is the additive inverse of the evidence lower bound objective (ELBO); in this work, we will refer to this as the ELBO loss, denoted $\mathcal{L}_{\text{ELBO}}$. By the non-negativity of the KL divergence, it serves as a variational bound on the negative log-likelihood of the observed variable \mathbf{x} :

$$-\log p_{\theta}(\mathbf{x}) \le \mathcal{L}_{\text{ELBO}}(\mathbf{x}) \tag{2.24}$$

$$= \mathbb{E}_{\mathbf{z} \sim q_{\phi}(\mathbf{z}|\mathbf{x})} \left[-\log \left(\frac{p_{\theta}(\mathbf{x}, \mathbf{z})}{q(\mathbf{z}|\mathbf{x})} \right) \right]$$
 (2.25)

As such, minimisation of the ELBO loss accomplishes two things. Firstly, it will approximately minimise the negative log-likelihood of the observed variable \mathbf{x} —the overriding goal of a likelihood-based generative model. Secondly, it will minimise the KL divergence of $q_{\phi}(\mathbf{z}|\mathbf{x})$ from $p_{\theta}(\mathbf{z}|\mathbf{x})$, thus encouraging the approximate posterior $q_{\phi}(\mathbf{z}|\mathbf{x})$ to approximate the true posterior $p_{\theta}(\mathbf{z}|\mathbf{x})$ as closely as possible.

2.5.3 Markovian Hierarchical Variational Autoencoders

The Markovian hierarchical variational autoencoder (MHVAE) [9, 15] is a versatile extension of the VAE, accommodating an unrestricted number $N_z \ge 1$ of latent variables. Notably, the joint distribution of the observed variable and the latent variables is Markovian:

$$p_{\theta}(\mathbf{x}, \mathbf{z}_1, \dots, \mathbf{z}_{N_z}) = p(\mathbf{x}|\mathbf{z}_1)p_{\theta}(\mathbf{z}_{N_z}) \prod_{i=1}^{N_z - 1} p_{\theta}(\mathbf{z}_i|\mathbf{z}_{i+1})$$
(2.26)

Figure 2.2 illustrates the MHVAE model. For the generative model, the observed variable \mathbf{x} is conditionally independent of \mathbf{z}_2 and \mathbf{z}_3 given \mathbf{z}_1 . Similarly, \mathbf{z}_1 is conditionally independent of \mathbf{z}_3 given \mathbf{z}_2 .

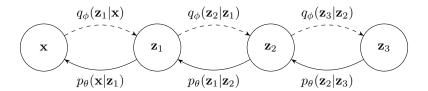


Figure 2.2: Graphical depiction of Markovian hierarchical VAE with one observed variable \mathbf{x} and three latent variables \mathbf{z}_1 , \mathbf{z}_2 and \mathbf{z}_3 . Solid lines depict the Bayesian network of the generative model; dashed lines depict the Bayesian network of the approximate inference model.

2.5.4 Infinitely Deep Markovian Hierarchical Variational Autoencoders

In the limit of $N_{\mathbf{z}} \to \infty$, we instead notationally write our latent variables in terms of a continuous-time variable $t \in [0, 1]$ as:

$$\{\mathbf{z}_0, \dots, \mathbf{z}_1\} = \{\mathbf{z}_t \mid t \in [0, 1]\}$$
 (2.27)

We can formulate the ELBO loss for the continuous-time MHVAE as follows:

$$-\log p_{\theta}(\mathbf{x}) \le \mathcal{L}_{\text{ELBO}}(\mathbf{x}) \tag{2.28}$$

$$= \mathbb{E}_{\mathbf{z}_0, \dots, \mathbf{z}_1 \sim q_{\phi}(\mathbf{z}_0, \dots, \mathbf{z}_1 | \mathbf{x})} \left[-\log \left(\frac{p_{\theta}(\mathbf{x}, \mathbf{z}_0, \dots, \mathbf{z}_1)}{q_{\phi}(\mathbf{z}_0, \dots, \mathbf{z}_1 | \mathbf{x})} \right) \right]$$
(2.29)

$$= \mathbb{E}_{\mathbf{z}_0 \sim q(\mathbf{z}_0, \mathbf{x})} \left[-\log p_{\theta}(\mathbf{x} | \mathbf{z}_0) \right] + D_{KL}(q(\mathbf{z}_0, \dots, \mathbf{z}_1 | \mathbf{x}) || p_{\theta}(\mathbf{z}_0, \dots, \mathbf{z}_1))$$
(2.30)

Per datapoint \mathbf{x} , we define $\mathcal{L}_T(t)$ as the KL divergence of $q_{\phi}(\mathbf{z}_t, \dots, \mathbf{z}_1 | \mathbf{x})$ from $p_{\theta}(\mathbf{z}_t, \dots, \mathbf{z}_1)$ for a subset of timesteps from t to 1, and its corresponding time derivative $\mathcal{L}'_T(t)$ as:

$$\mathcal{L}_T(t) = D_{KL}(q_{\phi}(\mathbf{z}_t, \dots, \mathbf{z}_1 | \mathbf{x}) || p_{\theta}(\mathbf{z}_t, \dots, \mathbf{z}_1))$$
(2.31)

$$\mathcal{L}_T'(t) = \frac{d}{dt}\mathcal{L}_T(t) \tag{2.32}$$

We can substitute this into the ELBO loss and use the second fundamental theorem of calculus to yield the following form for the ELBO loss, defined per datapoint \mathbf{x} as:

$$\mathcal{L}_{\text{ELBO}}(\mathbf{x}) = \mathbb{E}_{\mathbf{z}_0 \sim q(\mathbf{z}_0, \mathbf{x})} \left[-\log p_{\theta}(\mathbf{x} | \mathbf{z}_0) \right] + D_{KL}(q(\mathbf{z}_0, \dots, \mathbf{z}_1 | \mathbf{x}) || p_{\theta}(\mathbf{z}_0, \dots, \mathbf{z}_1))$$
(2.33)

$$= \mathbb{E}_{\mathbf{z}_0 \sim q(\mathbf{z}_0, \mathbf{x})} \left[-\log p_{\theta}(\mathbf{x} | \mathbf{z}_0) \right] + \mathcal{L}_T(0)$$
(2.34)

$$= \mathbb{E}_{\mathbf{z}_0 \sim q(\mathbf{z}_0, \mathbf{x})} \left[-\log p_{\theta}(\mathbf{x} | \mathbf{z}_0) \right] + \mathcal{L}_T(1) - \int_0^1 \mathcal{L}_T'(t) dt$$
 (2.35)

This form for the ELBO may seem unconventional. However, we introduce it here to motivate a strong theoretical link between the ELBO and the weighted loss [7], which is used to train diffusion models in the broader literature. We explore the weighted loss in Section 2.7.8.

2.6 Score-Based Generative Models

Score-based generative models [16, 17] are an alternative to likelihood-based generative models. We define the *score* of a probability density $p(\mathbf{x})$ to be:

$$\nabla_{\mathbf{x}} \log p(\mathbf{x}) \tag{2.36}$$

The score network \mathbf{s}_{θ} is a neural network parameterised by θ trained to approximate the score of the true data distribution $p^*(\mathbf{x})$, such that for any observed \mathbf{x} :

$$\mathbf{s}_{\theta}(\mathbf{x}) \approx \nabla_{\mathbf{x}} \log p^*(\mathbf{x})$$
 (2.37)

Importantly, we accomplish this without training a model to directly approximate the true data distribution $p^*(\mathbf{x})$ in advance.

2.7 Diffusion Models

2.7.1 Overview

Diffusion models [14, 3, 17] are a framework for generative modelling. Diffusion models consist of a forward diffusion process that transforms a datapoint into noise, and a reverse-time generative model able to transform noise back into a datapoint. As we explore in this section, diffusion models have both likelihood and score-based interpretations.

2.7.2 Forward Diffusion Process

Specified in continuous time, the *forward diffusion process* is a Gaussian diffusion process that defines the model's latent variables as a sequence of increasingly noisy versions of \mathbf{x} :

$$\{\mathbf{z}_0, \dots, \mathbf{z}_1\} = \{\mathbf{z}_t \mid t \in [0, 1]\}$$
 (2.38)

An Itô stochastic differential equation (SDE) defines the time evolution of the diffusion process [17]:

$$d\mathbf{z}_t = \mathbf{f}(\mathbf{z}_t, t)dt + g(t)d\mathbf{w}_t \tag{2.39}$$

where \mathbf{w}_t is the standard Wiener process (i.e. Brownian motion); $\mathbf{f}(\mathbf{z}_t, t) : \mathbb{R}^D \to \mathbb{R}^D$ is a vector-valued function called the *drift* coefficient of \mathbf{z} ; $g(t) : \mathbb{R} \to \mathbb{R}$ is a scalar-function known as the *diffusion* coefficient of \mathbf{z}_t ; and D is the dimensionality of our input data. In this work, we use a *variance-preserving* diffusion model, which we define by the following drift and diffusion coefficients:

$$\mathbf{f}(\mathbf{z}_t, t) = -\frac{1}{2} \left(\frac{d}{dt} \log \left(1 + \exp(-\lambda_t) \right) \right) \mathbf{z}_t$$
 (2.40)

$$g(t)^{2} = \frac{d}{dt}\log\left(1 + \exp(-\lambda_{t})\right) \tag{2.41}$$

where $\lambda_t \in [\lambda_{\min}, \lambda_{\max}]$ is a monotonically decreasing scalar-valued function of time $t \in [0, 1]$; we provide more details on λ_t in Section 2.7.4. The forward process forms a conditional joint distribution $q(\mathbf{z}_0, \dots, \mathbf{z}_1 | \mathbf{x})$, whose marginal distribution of each latent variable \mathbf{z}_t given the observed variable \mathbf{x} is given by:

$$q(\mathbf{z}_t|\mathbf{x}) = \mathcal{N}\left(\mathbf{z}_t; \alpha_t \mathbf{x}, \sigma_t^2 \mathbf{I}\right)$$
(2.42)

where α_t and σ_t are functions of λ_t , such that:

$$\alpha_t^2 = S(\lambda_t) \tag{2.43}$$

$$\sigma_t^2 = S(-\lambda_t) \tag{2.44}$$

where S is the sigmoid function. The joint distribution of latent variables $\mathbf{z}_r, \mathbf{z}_s, \mathbf{z}_t$ at subsequent timesteps $0 \le r < s < t \le 1$ is Markovian, and thus:

$$q(\mathbf{z}_t|\mathbf{z}_s, \mathbf{z}_r) = q(\mathbf{z}_t|\mathbf{z}_s) = \mathcal{N}\left(\mathbf{z}_t; \alpha_{t|s}\mathbf{z}_s, \sigma_{t|s}^2 \mathbf{I}\right)$$
(2.45)

where $\alpha_{t|s}$ and $\sigma_{t|s}$ are given by:

$$\alpha_{t|s} = \frac{\alpha_t}{\alpha_s} \tag{2.46}$$

$$\sigma_{t|s}^2 = \sigma_t^2 - \alpha_{t|s}^2 \sigma_s^2 \tag{2.47}$$

A full derivation of $q(\mathbf{z}_t|\mathbf{z}_s)$ is given in Appendix A.1. The true distribution $p^*(\mathbf{x})$ plus the conditional joint distribution of the forward model $q(\mathbf{z}_0, \dots, \mathbf{z}_1|\mathbf{x})$ defines the following joint distribution:

$$q(\mathbf{z}_0, \dots, \mathbf{z}_1) = \int p^*(\mathbf{x}) q(\mathbf{z}_0, \dots, \mathbf{z}_1 | \mathbf{x}) d\mathbf{x}$$
 (2.48)

2.7.3 Reverse-Time Generative Model

Anderson [1] shows that the forward SDE of Equation 2.39 is exactly solved by a second diffusion process, running backwards in time and given by the reverse-time SDE:

$$d\mathbf{z}_t = \left[\mathbf{f}(\mathbf{z}_t, t) - g(t)^2 \nabla_{\mathbf{z}_t} \log q(\mathbf{z}_t) \right] dt + g(t) d\bar{\mathbf{w}}_t$$
 (2.49)

where $\bar{\mathbf{w}}_t$ is a standard Wiener process when time flows backwards. Let $\mathbf{s}_{\theta}(\mathbf{z}_t, \lambda_t)$ be a λ_t -dependent score network [16] that approximates the score of $q(\mathbf{z}_t)$ such that:

$$\mathbf{s}_{\theta}(\mathbf{z}_{t}, \lambda_{t}) \approx \nabla_{\mathbf{z}_{t}} \log q(\mathbf{z}_{t})$$
 (2.50)

Thus, if we have a perfect score model $\mathbf{s}_{\theta}(\mathbf{z}_{t}, \lambda_{t}) = \nabla_{\mathbf{z}_{t}} q(\mathbf{z}_{t})$, then the reverse-time SDE is exactly:

$$d\mathbf{z}_{t} = \left[\mathbf{f}(\mathbf{z}_{t}, t) - g(t)^{2} \mathbf{s}_{\theta}(\mathbf{z}_{t}, \lambda_{t}) \right] dt + g(t) d\bar{\mathbf{w}}_{t}$$
(2.51)

In the broader literature, diffusion models utilise a variety of numerical solvers to provide approximate trajectories of the reverse-time SDE. In this work, we sequentially generate latent variables starting from t=1 and working backwards to t=0, over T uniformly-spaced discrete timesteps. More formally, this comprises a hierarchical generative model that defines a joint distribution over latent variables as follows:

$$p_{\theta}(\mathbf{z}_0, \dots, \mathbf{z}_1) = p_{\theta}(\mathbf{z}_1) \prod_{i=1}^{T} p_{\theta}(\mathbf{z}_{s(i)} | \mathbf{z}_{t(i)})$$
(2.52)

where $s(i) = (i-1) \cdot T^{-1}$ and $t(i) = i \cdot T^{-1}$. For large enough λ_{max} , \mathbf{z}_0 is almost noiseless, so learning a model $p_{\theta}(\mathbf{z}_0)$ is practically equivalent to learning a model $p_{\theta}(\mathbf{x})$. For sufficiently small λ_{min} , \mathbf{z}_1 contains almost no information about \mathbf{x} . As such, there exists a distribution $p_{\theta}(\mathbf{z}_1)$ such that:

$$D_{KL}(q(\mathbf{z}_1|\mathbf{x})||p_{\theta}(\mathbf{z}_1)) \approx 0 \tag{2.53}$$

For variance-preserving diffusion models, as used in this work, we model $p_{\theta}(\mathbf{z}_1)$ as the multivariate standard Gaussian:

$$p_{\theta}(\mathbf{z}_1) = \mathcal{N}(\mathbf{z}_1; \mathbf{0}, \mathbf{I}) \tag{2.54}$$

Once we have sampled $\mathbf{z}_1 \sim p_{\theta}(\mathbf{z}_1)$, we use the discrete-time ancestral sampler [3] to sequentially generate each latent variable \mathbf{z}_s from \mathbf{z}_t where $0 \le s < t \le 1$. This corresponds to a particular discretisation of the reverse-time variance-preserving SDE, as shown by Song et al. [17]. More formally, from a given latent \mathbf{z}_t we generate $\mathbf{z}_s \sim p_{\theta}(\mathbf{z}_s|\mathbf{z}_t)$ via:

$$p_{\theta}(\mathbf{z}_s|\mathbf{z}_t) = q(\mathbf{z}_s|\mathbf{z}_t, \mathbf{x} = \hat{\mathbf{x}}_{\theta}(\mathbf{z}_t, \lambda_t))$$
(2.55)

$$= \mathcal{N}\left(\tilde{\boldsymbol{\mu}}_{s|t}(\mathbf{z}_t, \mathbf{x} = \hat{\mathbf{x}}_{\theta}(\mathbf{z}_t, \lambda_t)), \tilde{\sigma}_{s|t}\mathbf{I}\right)$$
(2.56)

where $\hat{\mathbf{x}}_{\theta}(\mathbf{z}_t, \lambda_t)$ is our denoised estimate of the original data \mathbf{x} given latent \mathbf{z}_t and log signal-to-noise ratio λ_t , and

$$\tilde{\boldsymbol{\mu}}_{s|t}(\mathbf{z}_t, \mathbf{x}) = \frac{\alpha_{t|s}\sigma_s^2}{\sigma_t^2} \mathbf{z}_t + \frac{\alpha_s \sigma_{t|s}^2}{\sigma_t^2} \mathbf{x}$$
(2.57)

$$\tilde{\sigma}_{s|t} = \frac{\sigma_{t|s}\sigma_s}{\sigma_t} \tag{2.58}$$

Interestingly, in Equation 2.55 we introduced a denoiser network $\hat{\mathbf{x}}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$ to define the discrete-time ancestral sampler, while in Equation 2.50 we described diffusion models as learning a score network $\mathbf{s}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$. One of the powerful aspects of diffusion models is that we can freely switch between different parameterisations. For example, we can train a neural network $\mathbf{s}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$ to predict the score of \mathbf{z}_{t} and then convert the output to a denoised estimate of \mathbf{z}_{t} , as if we had trained a denoiser network $\mathbf{x}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$ directly. This reparameterisation property can be exceptionally advantageous. In this work, we explicitly train neither a score nor a denoiser network but rather use the \mathbf{v} -prediction parameterisation [13]; we describe this in detail in Section 2.7.5.

2.7.4 Noise Schedule

We formalise the notion that \mathbf{z}_t is increasingly noisy by defining the log signal-to-noise ratio

$$\lambda_t = \log\left(\frac{\alpha_t^2}{\sigma_t^2}\right) \in [\lambda_{\min}, \lambda_{\max}] \tag{2.59}$$

as a strictly monotonically decreasing function f_{Λ} of time $t \in [0,1]$, known as the noise schedule.

In this work, we use a truncated continuous-time version of the α -cosine schedule [10], introduced in its original discrete-time form by Nichol and Dhariwal [10]. The α -cosine schedule was motivated by the fact that the 'linear' schedule introduced in prior work by Ho et al. [3] causes α_t to fall to zero more quickly than is optimal. Nichol and Dhariwal empirically found that this induces too much noise in the latter stages of the forward diffusion process; as such, the latent variables \mathbf{z}_t in these stages contribute little to sample quality. In response, they proposed the original discrete-time α -cosine schedule. In this work, we use a continuous-time diffusion model and therefore use an adapted model described in [6]. More formally, we define:

$$\lambda_t = f_{\Lambda}(t) = -2\log\left(\tan\left(\frac{\pi}{2}(t_0 + t(t_1 - t_0))\right)\right)$$
 (2.60)

where t_0 and t_1 truncate $f_{\Lambda}(t)$ to the desired range $[\lambda_{\min}, \lambda_{\max}]$ for $t \in [0, 1]$, and are themselves defined as:

$$t_0 = \frac{2}{\pi} \arctan\left(\exp\left(-\frac{1}{2}\lambda_{\max}\right)\right) \tag{2.61}$$

$$t_1 = \frac{2}{\pi} \arctan\left(\exp\left(-\frac{1}{2}\lambda_{\min}\right)\right) \tag{2.62}$$

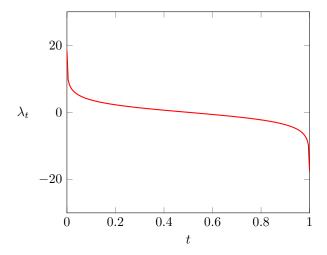


Figure 2.3: Relationship between time t and the log signal-to-noise ratio λ_t for the truncated continuoustime α -cosine noise schedule $f_{\Lambda}(t)$ as defined in Equation 2.60 with $\lambda_{\min} = -30$ and $\lambda_{\max} = 30$. The horizontal axis is time $t \in [0, 1]$; the vertical axis is $\lambda_t = f_{\Lambda}(t) \in [\lambda_{\min}, \lambda_{\max}] = [-30, 30]$.

Figure 2.3 visualises how the log signal-to-noise ratio $\lambda_t \in [\lambda_{\min}, \lambda_{\max}]$ varies with time $t \in [0, 1]$ using the α -cosine schedule detailed above.

We can compute α_t and σ_t from either λ_t or t via the following equations:

$$\alpha_t = \sqrt{S(\lambda_t)} = \cos\left(\frac{\pi}{2}(t_0 + t(t_1 - t_0))\right) \tag{2.63}$$

$$\alpha_t = \sqrt{S(\lambda_t)} = \cos\left(\frac{\pi}{2}(t_0 + t(t_1 - t_0))\right)$$

$$\sigma_t = \sqrt{S(-\lambda_t)} = \sin\left(\frac{\pi}{2}(t_0 + t(t_1 - t_0))\right)$$
(2.63)

where S is the sigmoid function. Figure 2.4 visualises how the values of α_t and σ_t vary with time $t \in [0,1]$ using the α -cosine schedule detailed above. Appendix A.2 provides further details on the form of f_{Λ} and how we can derive the forms for α_t and σ_t .

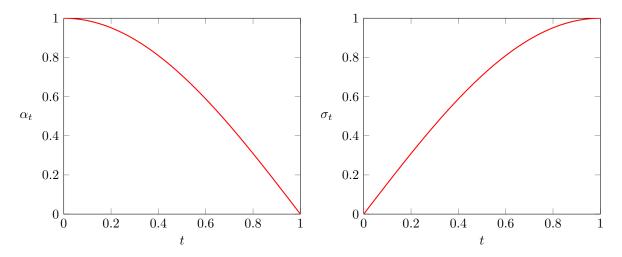


Figure 2.4: Relationship between time t and α_t (left) and σ_t (right) for the same truncated continuoustime α -cosine noise schedule as that in Figure 2.3. The horizontal axis is time $t \in [0,1]$; the vertical axis is the value of α_t (left) and σ_t (right).

We can, in theory, use two different noise schedules: one to train the model and another to generate new samples. During training, the noise schedule affects the variance of the gradients [7]. During generation, we typically want to use a noise schedule that optimises the quality of the generated samples. However, in this work, we use the truncated continuous-time α -cosine schedule for training and generation, as we found it to provide empirically good results for both. During training, we sampling $t \sim \mathcal{U}(0,1)$ uniformly at random, then compute $\lambda = f_{\Lambda}(t)$, which equates to sampling $\lambda \sim p_{\Lambda}(\lambda)$, where $p_{\Lambda}(\lambda)$ is the probability density function for the truncated continuous-time α -cosine schedule, and given by:

$$p_{\Lambda}(\lambda) = \frac{1}{2\pi(t_1 - t_0)} \operatorname{sech}\left(\frac{\lambda}{2}\right)$$
 (2.65)

Figure 2.5 displays the probability density function.

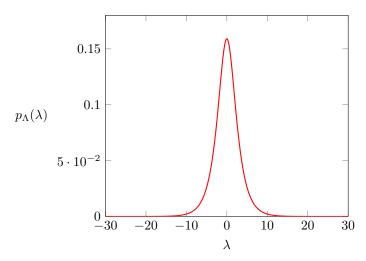


Figure 2.5: Probability density function $p_{\Lambda}(\lambda)$ for the same truncated continuous-time α -cosine schedule as that in Figure 2.3. The horizontal axis is the log signal-to-noise ratio $\lambda \in [\lambda_{\min}, \lambda_{\max}]$; the vertical axis is the corresponding probability density p_{Λ} .

2.7.5 Parameterisations

In Section 2.7.3, we defined our generative model $p_{\theta}(\mathbf{x})$ using $\hat{\mathbf{x}}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$, which takes as input some noisy latent variable \mathbf{z}_{t} and a log signal-to-noise ratio λ_{t} and outputs a denoised estimate of the latent. Training a neural network to predict $\mathbf{x} \approx \hat{\mathbf{x}}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$ directly is referred to as the \mathbf{x} -prediction parameterisation, but is seldom adopted in the broader literature due to sub-optimal results [3]. Recent diffusion models have instead adopted different parameterisations, most commonly the ϵ -prediction parameterisation (see e.g. [3, 4, 12]), wherein a neural network is instead trained to predict the noise $\epsilon \approx \hat{\epsilon}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$, from which we can compute a denoised estimate of noisy latent \mathbf{z}_{t} via:

$$\hat{\mathbf{x}}_{\theta}(\mathbf{z}_{t}, \lambda_{t}) = \frac{1}{\alpha_{t}} \left(\mathbf{z}_{t} - \sigma_{t} \hat{\boldsymbol{\epsilon}}_{\theta}(\mathbf{z}_{t}, \lambda_{t}) \right)$$
(2.66)

In this work, we employ the **v**-prediction parameterisation, introduced originally by Salimans and Ho [13], and commonly employed in video diffusion models (see e.g. [5, 2]). The **v**-prediction parameterisation was introduced initially to facilitate progressive distillation for faster sampling, though we utilise it here for its additional benefits highlighted by Ho et al. [2], namely faster convergence of sample quality and prevention of temporal colour shifting sometimes observed with ϵ -prediction video diffusion models.

Formally, for a given datapoint $\mathbf{x} \sim q(\mathbf{x})$ we define the velocity of $\mathbf{z}_t \sim q(\mathbf{z}_t|\mathbf{x})$ as:

$$\mathbf{v}_t = \alpha_t \boldsymbol{\epsilon} - \sigma_t \mathbf{x} \tag{2.67}$$

where $\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$ is multivariate standard Gaussian noise. We train our neural network $\hat{\mathbf{v}}_{\theta}(\mathbf{z}_t, \lambda_t)$ to minimise the following loss function, defined per datapoint \mathbf{x} as:

$$\mathbb{E}_{\lambda \sim p_{\Lambda}(\lambda), \epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})} \left[\| \mathbf{v}_t - \hat{\mathbf{v}}_{\theta}(\mathbf{z}_t, \lambda_t) \|_2^2 \right]$$
 (2.68)

During discrete-time ancestral sampling, we convert our estimate $\mathbf{v}_t \approx \hat{\mathbf{v}}_{\theta}(\mathbf{z}_t, \lambda_t)$ into an estimate of the denoised latent $\mathbf{x} \approx \hat{\mathbf{x}}_{\theta}(\mathbf{z}_t, \lambda_t)$ via:

$$\hat{\mathbf{x}}(\mathbf{z}_t, \lambda_t) = \alpha_t \mathbf{z}_t - \sigma_t \hat{\mathbf{v}}_\theta(\mathbf{z}_t, \lambda_t) \tag{2.69}$$

Appendix A.3 provides further details on the **v**-prediction parameterisation, including derivations of the velocity and denoised latent.

2.7.6 Score-Based Interpretation

Suppose we have a multivariate Gaussian variable with mean μ and covariance Σ :

$$\mathbf{z} \sim p(\mathbf{z}) = \mathcal{N}(\mathbf{z}; \boldsymbol{\mu}_{\mathbf{z}}, \boldsymbol{\Sigma}_{\mathbf{z}})$$
 (2.70)

Tweedie's formula states that:

$$\mathbb{E}\left[\boldsymbol{\mu}_{\mathbf{z}}|\mathbf{z}\right] = \mathbf{z} + \boldsymbol{\Sigma}_{\mathbf{z}}\nabla_{\mathbf{z}}\log p(\mathbf{z}) \tag{2.71}$$

As such, for a given latent $\mathbf{z}_t \sim \mathcal{N}(\mathbf{z}_t; \alpha_t \mathbf{x}, \sigma_t^2 \mathbf{I})$, the expected value of the mean $\boldsymbol{\mu}_{\mathbf{z}_t}$ given \mathbf{z}_t is given by:

$$\mathbb{E}\left[\boldsymbol{\mu}_{\mathbf{z}_t}|\mathbf{z}_t\right] = \mathbf{z}_t + \sigma_t^2 \nabla_{\mathbf{z}_t} \log q(\mathbf{z}_t)$$
(2.72)

From Equation 2.42, we have $\mu_{\mathbf{z}_t} = \alpha_t \mathbf{x}$. Thus, we can reformulate the score of $q(\mathbf{z}_t)$ in terms of \mathbf{z}_t and \mathbf{x} as:

$$\alpha_t \mathbf{x} = \mathbf{z}_t + \sigma_t^2 \nabla_{\mathbf{z}_t} \log q(\mathbf{z}_t) \tag{2.73}$$

2.7.7 ELBO for Diffusion Models

We can interpret the variance-preserving diffusion model used in this work as an MHVAE with several additional restrictions. Firstly, the dimensionality of each latent \mathbf{z}_t equals the dimensionality of the observed variable. Secondly, we have pre-defined $q(\mathbf{z}_t|\mathbf{z}_s)$ where $0 \le s < t \le 1$ as a Gaussian diffusion process with no learnable inference parameters. Finally, the marginal distribution of the final latent $q(\mathbf{z}_1)$ is approximately the multivariate standard Gaussian $\mathcal{N}(\mathbf{0}, \mathbf{I})$, and thus holds effectively no information about the observed variable \mathbf{x} . VAEs and MHVAEs do not typically have these restrictions. Nonetheless, much like VAEs and MHVAEs, we can optimise the generative parameters θ of diffusion models by minimising the ELBO loss. As a notable example, Sohl-Dickstein et al. [14] optimised the original discrete-time diffusion model via the ELBO loss.

For a given datapoint \mathbf{x} , we define $\mathcal{L}_{\Lambda}(\lambda)$ as the KL divergence of $q(\mathbf{z}_t, \dots, \mathbf{z}_1 | \mathbf{x})$ from $p_{\theta}(\mathbf{z}_t, \dots, \mathbf{z}_1)$ for a subset of timesteps from $t = f_{\Lambda}^{-1}(\lambda)$ to 1 for datapoint \mathbf{x} :

$$\mathcal{L}_{\Lambda}(\lambda) = D_{KL}(q(\mathbf{z}_t, \dots, \mathbf{z}_1 | \mathbf{x}) || p_{\theta}(\mathbf{z}_t, \dots, \mathbf{z}_1))$$
(2.74)

Notably, $\mathcal{L}_{\Lambda}(\lambda)$ equates to $\mathcal{L}_{T}(t)$ defined in Equation 2.31 under a simple change of variable:

$$\mathcal{L}_{\Lambda}(\lambda) = \mathcal{L}_{T}(t = f_{\Lambda}^{-1}(\lambda)) \tag{2.75}$$

Similarly, we can reformulate the ELBO loss for a continuous-time MHVAE given in Equation 2.35 to provide the ELBO loss in terms of the log signal-to-noise ratio λ ; it is given per datapoint \mathbf{x} by:

$$\mathcal{L}_{\text{ELBO}}(\mathbf{x}) = \mathbb{E}_{\mathbf{z}_0 \sim q(\mathbf{z}_0, \mathbf{x})} \left[-\log p_{\theta}(\mathbf{x} | \mathbf{z}_0) \right] + \mathcal{L}_{\Lambda}(\lambda_{\text{max}})$$
(2.76)

$$= \underbrace{\mathbb{E}_{\mathbf{z}_0 \sim q(\mathbf{z}_0, \mathbf{x})} \left[-\log p_{\theta}(\mathbf{x} | \mathbf{z}_0) \right]}_{\text{Reconstruction Loss}} + \underbrace{\mathcal{L}_{\Lambda}(\lambda_{\min})}_{\text{Prior Loss}} + \int_{\lambda_{\min}}^{\lambda_{\max}} \mathcal{L}'_{\Lambda}(\lambda) d\lambda$$
(2.77)

With sufficiently large λ_{max} , the reconstruction loss is approximately zero since we can almost perfectly reconstruct \mathbf{x} from \mathbf{z}_0 —this is particularly true for discrete \mathbf{x} . Mathematically, as $\lambda_{\text{max}} \to \infty$, we have:

$$\lim_{\lambda_{\text{max}} \to \infty} q(\mathbf{z}_0 | \mathbf{x}) = \delta(\mathbf{z}_0 - \mathbf{x})$$
(2.78)

where δ is the Dirac delta distribution. Similarly, with sufficiently small λ_{\min} , the prior loss is approximately zero; as $\lambda_{\min} \to -\infty$, we have:

$$\lim_{\lambda_{\min} \to -\infty} q(\mathbf{z}_1 | \mathbf{x}) = \mathcal{N}(\mathbf{0}, \mathbf{I}) = p_{\theta}(\mathbf{z}_1)$$
(2.79)

so the KL divergence prior loss term likewise approaches zero.

In Appendix C of [7], Kingma and Gao showed that $\mathcal{L}'_{\Lambda}(\lambda)$ —which, with a slight abuse of terminology, they refer to as the *time derivative*—simplifies to a remarkable degree:

$$\mathcal{L}'_{\Lambda}(\lambda) = \frac{d}{d\lambda} \mathcal{L}_{\Lambda}(\lambda) = \frac{1}{2} \mathbb{E}_{\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})} \left[\| \epsilon - \hat{\epsilon}_{\theta}(\mathbf{z}_{t}, \lambda_{t}) \|_{2}^{2} \right]$$
(2.80)

2.7.8 Weighted Loss

Most diffusion models in the broader literature—including state-of-the-art models—do not optimise their parameters θ via minimisation of the ELBO loss. In practice, the various objectives used are all special cases of a *weighted loss* [7], which is defined per datapoint \mathbf{x} as:

$$\mathcal{L}_{\text{WL}} = w(\lambda_{\min}) \mathcal{L}_{\Lambda}(\lambda_{\min}) + \int_{\lambda_{\min}}^{\lambda_{\max}} w(\lambda) \mathcal{L}'_{\Lambda}(\lambda) d\lambda$$
 (2.81)

where $w(\lambda)$ is a weighting function. Note that, assuming the reconstruction loss is approximately zero, the ELBO loss given in Equation 2.77 is a special case of the weighted loss \mathcal{L}_{WL} with $w(\lambda) = 1$. Substituting the form of $\mathcal{L}'_{\Lambda}(\lambda)$ given in Equation 2.80 yields the following form for the weighted loss:

$$\mathcal{L}_{\text{WL}} = w(\lambda_{\min}) \mathcal{L}_{\Lambda}(\lambda_{\min}) + \frac{1}{2} \int_{\lambda_{\min}}^{\lambda_{\max}} w(\lambda) \mathbb{E}_{\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})} \left[\| \epsilon - \hat{\epsilon}_{\theta}(\mathbf{z}_{t}, \lambda) \|_{2}^{2} \right] d\lambda$$
 (2.82)

This form provides several useful insights. Since the first term—the weighted prior loss—contains no learnable parameters, minimisation of the weighted loss \mathcal{L}_{WL} equates to minimisation of the intractable integral. In practice, we minimise the integral via an importance-weighted Monte Carlo integrator:

$$\int_{\lambda_{\min}}^{\lambda_{\max}} w(\lambda) \mathbb{E}_{\boldsymbol{\epsilon} \sim \mathcal{N}(\mathbf{0}, \mathbf{I})} \left[\| \boldsymbol{\epsilon} - \hat{\boldsymbol{\epsilon}}_{\theta}(\mathbf{z}_{t}, \lambda_{t}) \|_{2}^{2} \right] d\lambda = \mathbb{E}_{\lambda \sim p_{\Lambda}(\lambda)} \left[\frac{w(\lambda)}{p_{\Lambda}(\lambda)} \mathbb{E}_{\boldsymbol{\epsilon} \sim \mathcal{N}(\mathbf{0}, \mathbf{I})} \left[\| \boldsymbol{\epsilon} - \hat{\boldsymbol{\epsilon}}_{\theta}(\mathbf{z}_{t}, \lambda) \|_{2}^{2} \right] \right]$$
(2.83)

$$= \mathbb{E}_{\lambda \sim p_{\Lambda}(\lambda), \epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})} \left[\frac{w(\lambda)}{p_{\Lambda}(\lambda)} \| \epsilon - \hat{\epsilon}_{\theta}(\mathbf{z}_{t}, \lambda) \|_{2}^{2} \right]$$
(2.84)

$$\simeq \frac{w(\lambda)}{p_{\Lambda}(\lambda)} \| \boldsymbol{\epsilon} - \hat{\boldsymbol{\epsilon}}_{\theta}(\mathbf{z}_{t}, \lambda) \|_{2}^{2}$$
 (2.85)

Notable further analysis by Kingma and Gao [7] shows that the weighted loss also has a likelihood-based interpretation. Simple integration by parts enables us to write the weighted loss as:

$$\mathcal{L}_{\text{WL}} = w(\lambda_{\text{max}}) \mathcal{L}_{\Lambda}(\lambda_{\text{max}}) + \int_{\lambda_{\text{min}}}^{\lambda_{\text{max}}} -w'(\lambda) \mathcal{L}_{\Lambda}(\lambda) d\lambda$$
 (2.86)

The likelihood-based interpretation comes from the fact that $\mathcal{L}_{\Lambda}(\lambda)$ serves as a variational bound on the negative marginal likelihood of the noise-perturbed data $\mathbf{z}_t \sim q(\mathbf{z}_t|\mathbf{x})$:

$$\mathcal{L}_{\Lambda}(\lambda) \ge D_{KL}(q(\mathbf{z}_t|\mathbf{x})||p(\mathbf{z}_t)) \tag{2.87}$$

$$= \mathbb{E}_{\mathbf{z}_{t} \sim q(\mathbf{z}_{t}|\mathbf{x})} \left[-\log p(\mathbf{z}_{t}) \right] + \mathbb{E}_{\mathbf{z}_{t} \sim q(\mathbf{z}_{t}|\mathbf{x})} \left[\log q(\mathbf{z}_{t}|\mathbf{x}) \right]$$
(2.88)

As such, minimisation of $\mathcal{L}_{\Lambda}(\lambda)$ equates to maximisation of the expected log-likelihood of the noise-perturbed data $\mathbf{z}_t \sim q(\mathbf{z}_t|\mathbf{x})$ with noise level λ . If the weighting function $w(\lambda)$ is a monotonically decreasing function of $\lambda \in [\lambda_{\min}, \lambda_{\max}]$, then by definition $-w'(\lambda)$ will be positive for all $\lambda \in [\lambda_{\min}, \lambda_{\max}]$. In which case, minimisation of \mathcal{L}_{WL} itself equates to maximisation of the weighted expected log-likelihood of the noise-perturbed data $\mathbf{z}_t \sim q(\mathbf{z}_t|\mathbf{x})$ with weighting $-w'(\lambda)$.

Kingma and Gao's [7] analysis directly justifies our use of the **v**-prediction parameterisation with the truncated continuous-time α -cosine noise schedule. In conjunction, their use equates to the weighted loss with:

$$w(\lambda) = \frac{1}{2\pi(t_1 - t_0)} \exp\left(-\frac{\lambda}{2}\right) \tag{2.89}$$

$$-w'(\lambda) = \frac{1}{4\pi(t_1 - t_0)} \exp\left(-\frac{\lambda}{2}\right)$$
 (2.90)

Figure 2.6 shows the weighting function $w(\lambda)$ and the negative of its derivative $-w'(\lambda)$ for the v-parameterisation loss. As evident from the graphs, the weighting function $w(\lambda)$ is a monotonically decreasing function of λ , and as such $-w'(\lambda)$ is positive for all $\lambda \in [\lambda_{\min}, \lambda_{\max}]$. Therefore, during training, we are maximising a weighted expected log-likelihood of noise-perturbed data $\mathbf{z}_t \sim q(\mathbf{z}_t|\mathbf{x})$ for all $\lambda \in [\lambda_{\min}, \lambda_{\max}]$. In contrast, most diffusion models in the broader literature (see e.g. [3, 10, 12]) undergo training with non-monotonic weighting functions. In such cases, for noise levels whereby $-w'(\lambda)$ is negative, the weighted loss has a counterintuitive interpretation of minimisation of the weighted expected log-likelihood of the noise-perturbed data $\mathbf{z}_t \sim q(\mathbf{z}_t|\mathbf{x})$.

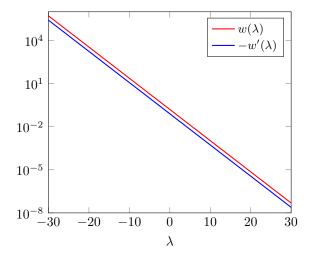


Figure 2.6: Relationship between the log signal-to-noise ratio λ and the functions $w(\lambda)$ and $-w'(\lambda)$ for the **v**-parameterisation with the same truncated continuous-time α -cosine schedule as that in Figure 2.3. The horizontal axis is the log signal-to-noise level λ ; the vertical axis is the value of $w(\lambda)$ and $-w'(\lambda)$; the vertical axis is logarithmic.

2.7.9 Imputation for Conditional Generation

In Section 2.2, we introduced the concept of conditional generation. In this work, however, we do not train a conditional model explicitly. Instead, we utilise reconstruction-guided sampling [5]: a sophisticated technique that derives a conditional model approximately from an unconditional model. More specifically, reconstruction-guided sampling facilitates the conditional generation of the unknown dimensions of some observed datapoint \mathbf{x} given the known dimensions. Deriving such a conditional model approximately from an unconditional model is advantageous: it enables us to train only a single unconditional model, which we can then flexibly use to facilitate the conditional generation of any unknown subset of dimensions. In this work, we utilise reconstruction-guided sampling to facilitate three distinct conditional generation tasks: temporal interpolation, forecasting, and autoregressive generation of arbitrarily long samples.

Reconstruction-guided sampling extends a prior technique known as *imputation*, introduced by Song et al. [17]. Thus, we first introduce imputation for completeness and to better motivate our use of reconstruction-guided sampling in this work.

We denote by $\Omega(\mathbf{x})$ and $\bar{\Omega}(\mathbf{x})$ the known and unknown dimensions of some observed datapoint \mathbf{x} , respectively. Formally, our goal is to derive the following conditional model without training it explicitly:

$$p_{\theta}(\bar{\Omega}(\mathbf{x})|\Omega(\mathbf{x})) \tag{2.91}$$

We can write the forward diffusion process for the unknown dimensions as the following SDE:

$$d\bar{\Omega}(\mathbf{z}_t) = \mathbf{f}(\bar{\Omega}(\mathbf{z}_t), t) + g(t)d\mathbf{w}_t \tag{2.92}$$

Anderson [1] shows that the corresponding reverse-time SDE conditioned on the known dimensions $\Omega(\mathbf{x})$ is given by:

$$d\bar{\Omega}(\mathbf{z}_t) = \left[\mathbf{f}(\bar{\Omega}(\mathbf{z}_t), t) - g(t)^2 \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t) | \Omega(\mathbf{x})) \right] dt + g(t) d\bar{\mathbf{w}}_t$$
 (2.93)

Although $q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x}))$ is intractable, we can approximate it as follows:

$$q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x})) = \int q(\bar{\Omega}(\mathbf{z}_t), \Omega(\mathbf{z}_t)|\Omega(\mathbf{x})) d\Omega(\mathbf{z}_t)$$
(2.94)

$$= \int q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x}), \Omega(\mathbf{z}_t))q(\Omega(\mathbf{z}_t)|\Omega(\mathbf{x}))d\Omega(\mathbf{z}_t)$$
(2.95)

$$= \mathbb{E}_{\Omega(\mathbf{z}_t) \sim q(\Omega(\mathbf{z}_t)|\Omega(\mathbf{x}))} \left[q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x}), \Omega(\mathbf{z}_t)) \right]$$
 (2.96)

$$\approx \mathbb{E}_{\Omega(\mathbf{z}_t) \sim q(\Omega(\mathbf{z}_t)|\Omega(\mathbf{x}))} \left[q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{z}_t)) \right]$$
 (2.97)

Song et al. [17] argue that the approximation in Equation 2.97 is appropriate since for small t, $\Omega(\mathbf{x})$ is almost the same as $\Omega(\mathbf{z}_t)$; and for larger t, $\Omega(\mathbf{x})$ is further away from $\bar{\Omega}(\mathbf{z}_t)$ in the Markov chain, and thus

has a smaller impact on $\bar{\Omega}(\mathbf{z}_t)$. Assuming the approximation holds, we can derive an unbiased estimator of $q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x}))$ as:

$$q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x})) \approx \mathbb{E}_{\Omega(\mathbf{z}_t) \sim q(\Omega(\mathbf{z}_t)|\Omega(\mathbf{x}))} \left[q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{z}_t)) \right] \simeq q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{z}_t))$$
(2.98)

The score of the natural logarithm of the unbiased estimator is given by:

$$\nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{z}_t)) = \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t),\Omega(\mathbf{z}_t)) - \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\Omega(\mathbf{z}_t))$$
(2.99)

$$= \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t), \Omega(\mathbf{z}_t)) \tag{2.100}$$

$$= \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q([\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)])$$
 (2.101)

where $[\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)]$ denotes a vector such that:

$$\Omega([\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)]) = \Omega(\mathbf{z}_t) \tag{2.102}$$

$$\bar{\Omega}([\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)]) = \bar{\Omega}(\mathbf{z}_t) \tag{2.103}$$

Thus, assuming the assumption given in Equation 2.97 holds, we can consequently approximate the reverse-time SDE conditioned on $\Omega(\mathbf{x})$ given in Equation 2.93 as follows:

$$d\bar{\Omega}(\mathbf{z}_t) = \left[\mathbf{f}(\bar{\Omega}(\mathbf{z}_t), t) - g(t)^2 \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q([\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)])\right] dt + g(t) d\bar{\mathbf{w}}_t$$
(2.104)

If we have a perfect score model, $\mathbf{s}_{\theta}(\mathbf{z}_t, \lambda_t) = \nabla_{\mathbf{z}_t} q(\mathbf{z}_t)$, then the reverse-time SDE is thus given by:

$$d\bar{\Omega}(\mathbf{z}_t) = \left[\mathbf{f}(\bar{\Omega}(\mathbf{z}_t), t) - g(t)^2 \mathbf{s}_{\theta}([\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)], \lambda_t) \right] dt + g(t) d\bar{\mathbf{w}}_t$$
 (2.105)

To provide a more intuitive link between Equation 2.105 and the generative procedure in Section 3.2, we also define the imputation method as a conditional denoiser $\hat{\mathbf{x}}_{\theta}^{C}(\mathbf{z}_{t}, \lambda_{t}, \Omega(\mathbf{x}))$, which takes the known dimensions of \mathbf{x} as input:

$$\hat{\mathbf{x}}_{\theta}^{\mathrm{C}}(\mathbf{z}_{t}, \lambda_{t}, \Omega(\mathbf{x})) = \bar{\Omega}\left(\hat{\mathbf{x}}_{\theta}([\bar{\Omega}(\mathbf{z}_{t}); \Omega(\mathbf{z}_{t})], \lambda_{t})\right)$$
(2.106)

Thus, the imputation method for conditional generation yields only a slight adjustment to the generative procedure defined in Section 2.7.3. Namely, at each step of the discrete-time ancestral sampler, we replace the dimensions of \mathbf{z}_t corresponding to the known dimensions of \mathbf{x} with an exact sample from the forward process: $\Omega(\mathbf{z}_t) \sim q(\Omega(\mathbf{z}_t)|\Omega(\mathbf{x}))$.

2.7.10 Reconstruction-Guided Sampling for Conditional Generation

Ho et al. [5] showed that the imputation process produces incoherent samples when applied to video diffusion models. Namely, although a given $\bar{\Omega}(\mathbf{x})$ will often appear reasonable in isolation, it will often not be coherent with $\Omega(\mathbf{x})$. This incoherency is likely because the assumption in Equation 2.97 does not hold for all $t \in [0,1]$. Avoiding the assumption, we instead construct an unbiased estimator for $q(\bar{\Omega}(\mathbf{z}_t|\Omega(\mathbf{x})))$ as:

$$q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x})) = \mathbb{E}_{\Omega(\mathbf{z}_t) \sim q(\Omega(\mathbf{z}_t)|\Omega(\mathbf{x}))} \left[q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x}), \Omega(\mathbf{z}_t)) \right] \simeq q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x}), \Omega(\mathbf{z}_t))$$
(2.107)

The score of the natural logarithm of the unbiased estimator is given by:

$$\nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t) | \Omega(\mathbf{x}), \Omega(\mathbf{z}_t)) = \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t), \Omega(\mathbf{x}), \Omega(\mathbf{z}_t)) - \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\Omega(\mathbf{z}_t), \Omega(\mathbf{x}))$$
(2.108)

$$= \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t), \Omega(\mathbf{x}), \Omega(\mathbf{z}_t))$$
(2.109)

$$= \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t), \Omega(\mathbf{z}_t)) + \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\Omega(\mathbf{x})|\bar{\Omega}(\mathbf{z}_t), \Omega(\mathbf{z}_t))$$
(2.110)

$$= \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q([\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)]) + \nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\Omega(\mathbf{x}) | [\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)]) \quad (2.111)$$

The second term in Equation 2.111 is missing in the imputation approach. Plugging in this missing term would make conditional sampling exact. However, since $q(\Omega(\mathbf{x})|[\bar{\Omega}(\mathbf{z}_t);\Omega(\mathbf{z}_t)])$ is not available in closed form, we must approximate it. Ho et al. [5] proposed to approximate it with a multivariate Gaussian of the form:

$$q(\Omega(\mathbf{x})|[\bar{\Omega}(\mathbf{z}_t);\Omega(\mathbf{z}_t)]) \approx \mathcal{N}\left(\Omega(\mathbf{x});\Omega\left(\hat{\mathbf{x}}_{\theta}\left([\bar{\Omega}(\mathbf{z}_t);\Omega(\mathbf{z}_t)],\lambda_t\right)\right),\left(\frac{\sigma_t^2}{\alpha_t^2}\right)\mathbf{I}\right)$$
(2.112)

Under this approximation, the second term of Equation 2.111 is thus itself approximated by:

$$\nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\Omega(\mathbf{x}) | [\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)]) \approx -\frac{\alpha_t^2}{2\sigma_t^2} \nabla_{\bar{\Omega}(\mathbf{z}_t)} \|\Omega(\mathbf{x}) - \Omega\left(\hat{\mathbf{x}}_{\theta}\left([\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)], \lambda_t\right)\right) \|_2^2$$
(2.113)

Ho et al. [5] interpret the inclusion of this term—absent in the imputation method—as a form of guidance based on the model's reconstruction of the conditioning data. They found empirically that—as with other forms of guidance—including a large weighting term $w_r > 1$ tends to improve sample quality further. They refer to this technique as reconstruction-guided sampling. Formally, assuming we have a perfect score model $\mathbf{s}_{\theta}(\mathbf{z}_t, \lambda_t) = \nabla_{\mathbf{z}_t} q(\mathbf{z}_t)$, reconstruction-guided sampling derives a conditional model from an unconditional model by approximating the score given in Equation 2.93 by:

$$\nabla_{\bar{\Omega}(\mathbf{z}_t)} \log q(\bar{\Omega}(\mathbf{z}_t)|\Omega(\mathbf{x})) \approx \mathbf{s}_{\theta}([\bar{\Omega}(\mathbf{z}_t);\Omega(\mathbf{z}_t)],\lambda_t)$$
(2.114)

$$-\frac{w_r \alpha_t^2}{2\sigma_t^2} \nabla_{\bar{\Omega}(\mathbf{z}_t)} \|\Omega(\mathbf{x}) - \Omega\left(\hat{\mathbf{x}}_{\theta}\left([\bar{\Omega}(\mathbf{z}_t); \Omega(\mathbf{z}_t)], \lambda_t\right)\right)\|_2^2$$
 (2.115)

Writing the reconstruction-guidance method as a conditional denoiser $\hat{\mathbf{x}}_{\theta}^{\mathrm{C}}(\mathbf{z}_{t}, \lambda_{t}, \Omega(\mathbf{x}))$ yields:

$$\hat{\mathbf{x}}_{\theta}^{\mathrm{C}}(\mathbf{z}_{t}, \lambda_{t}, \Omega(\mathbf{x})) = \bar{\Omega}\left(\hat{\mathbf{x}}_{\theta}([\bar{\Omega}(\mathbf{z}_{t}); \Omega(\mathbf{z}_{t})], \lambda_{t})\right) - \frac{w_{r}\alpha_{t}}{2} \nabla_{\bar{\Omega}(\mathbf{z}_{t})} \left\|\Omega(\mathbf{x}) - \Omega\left(\hat{\mathbf{x}}_{\theta}\left([\bar{\Omega}(\mathbf{z}_{t}); \Omega(\mathbf{z}_{t})]\right)\right)\right\|_{2}^{2}$$
(2.116)

2.7.11 U-Net

Chapter 3

Climate Background

- 3.1 Climate Simulations
- 3.2 Generative Models for Climate Simulations

Chapter 4

Experiments and Results

4.1 Dataset

4.1.1 UKCP18

4.1.2 Train and Test Sets

We scale each input data element to the range [-1,1] to ensure that our neural network operates on consistently scaled inputs during the reverse-time generative process, starting from the multivariate standard Gaussian prior.

4.2 Importance of the Transformation

4.2.1 No Transformation

One of the primary contributions of this work is a demonstration that a transformation of the input data is critical to the model's performance. To motivate this, we first provide results for a model trained without transforming the input data. Notably, the model produced samples with significant perceptible noise at the lower end of the precipitation scale. Figure 4.1 depicts an example of this. While the sample is temporally coherent, the noise is abundantly visible and detrimental to the quality of the sample.

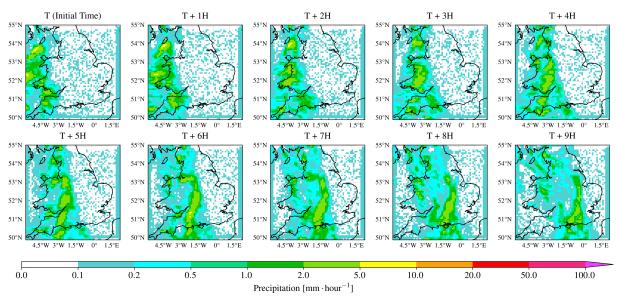


Figure 4.1: Randomly selected sample generated via our model trained with no transformation of the input data. The sample depicts hourly precipitation for the same $563.2~\mathrm{km} \times 563.2~\mathrm{km}$ region centred on Birmingham, UK, over ten hours. Each grid cell illustrates the mean precipitation rate, measured in millimetres per hour, for an $8.8~\mathrm{km} \times 8.8~\mathrm{km}$ subarea.

To motivate this beyond a single example, Figure 4.2 depicts the distribution of precipitation rates of our generated samples compared to those in our test set at the lower end of the scale (i.e. less than 0.5 millimetres per hour) where the noise is most perceptible. As can be seen, below approximately 0.2 millimetres per hour, the distribution of precipitation rates in our generated samples significantly deviates from that of the test set. For example, our generated samples contain precipitation rates of 0.1 millimetres per hour ten times more frequently than the test set. While 0.1 millimetres per hour equates to only very light drizzle, it is still a non-negligible amount of precipitation. The fact that the discrepancy is so large at this scale likely means that using the generated samples in any real-world context would present significant challenges.

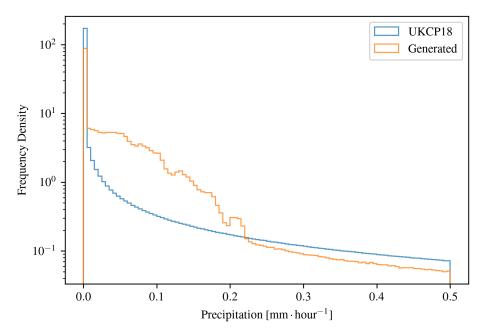


Figure 4.2: Distribution of precipitation rates below 0.5 millimetres per hour in samples generated via our model trained with no transformation compared to the test set from UKCP18. The horizontal axis is the precipitation rate, measured in millimetres per hour; the vertical axis is frequency density. The vertical axis is scaled logarithmically.

We can attribute much of this deviation to random noise visible in the generated samples. We cannot determine the precise source of this noise, but we speculate it may be a combination of multiple factors. The first contributing factor is likely that our score model $\mathbf{s}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$ is not perfect; in other words, there likely exists \mathbf{z}_{t} and λ_{t} such that $\mathbf{s}_{\theta}(\mathbf{z}_{t}, \lambda_{t})$ does not serve as a good approximation for $\nabla_{\mathbf{z}_{t}} \log q(\mathbf{z}_{t})$. In addition, since we can only approximate the reverse-time SDE, the discretised numerical solver likely introduces errors itself—this would be the case even if we had a perfect score model. We experimented with predictor-corrector sampling [17] to rectify this but observed no improvement in the quality of our generated samples. We speculate that the errors introduced by these two factors likely manifest as the noise in our generated samples.

Noise of this magnitude creates unique challenges for diffusion models generating precipitation data—challenges that do not present as significant an issue in the more common domain of image generation. There are two primary reasons for this. First and foremost, the distribution of precipitation rates in our data is far more skewed than the distribution of colours in image data. This skew is evident in Figure 4.3, which depicts the full distribution of precipitation rates in our test set. 80% of the precipitation rates are less than 0.003 millimetres per hour; to put this into perspective, the highest precipitation rate in our test set is approximately 60 millimetres per hour. As a direct result of this extreme distribution, even relatively small amounts of noise at the lower end of the scale can significantly impact the quality of the generated samples.

The second challenge that generating precipitation data presents in contrast to image data is that precipitation is a continuous variable. In contrast, discrete values comprise each colour channel in an image. Thus, a final discretisation step at the end of the generative process for images eliminates much of the noise. We cannot take advantage of this approach for precipitation data. In the context of Figure 4.1 wherein the noise is most perceptible below the 0.2 millimetres per hour threshold, this constitutes

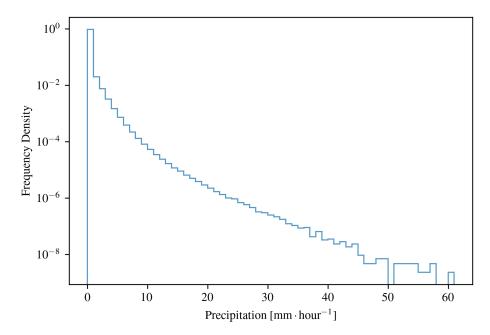


Figure 4.3: Distribution of precipitation rates in the test set from UKCP18. The horizontal axis is the precipitation rate, measured in millimetres per hour; the vertical axis is frequency density. The vertical axis is scaled logarithmically.

less than 0.2% of our output space. If this were an image with 256 discrete values per colour channel, this relative amount of noise represents less than a single discrete value. Thus, a final discretisation step would eliminate the noise almost entirely.

One final unnatural characteristic exhibited by a small proportion of the samples generated via our model trained with no transformation of the input data is an unnatural global shift in precipitation rates. Visually, this effect manifests as a consistent blue colour. Specifically, the samples in question exhibit a low precipitation rate in the region of 0.1--0.5 millimetres per hour across the entire sample, which represents a low drizzle rate across the entire $563.2 \text{ km} \times 563.2 \text{ km}$ region for the entire ten hours. Figure 4.4 depicts one such sample. Even in isolation, this would be an unnatural scenario and thus should certainly not occur at the frequency we observed in the samples we generated. We could not identify a single example in the test set that exhibited this characteristic.

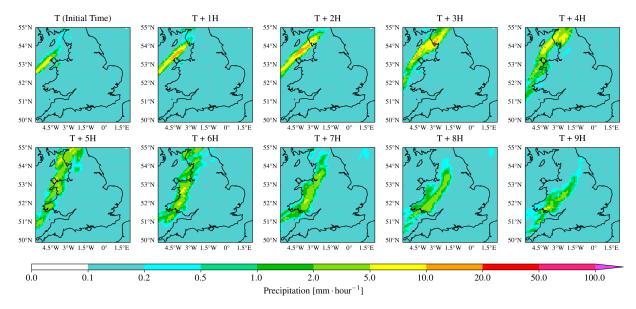


Figure 4.4: Sample generated via our model trained with no transformation of the input data that exhibits an unnatural global shift in precipitation rates. The format is the same as Figure 4.1.

4.2.2 Square Root Transformation

Empirically, we found that performing a square root transformation to the input data eliminated the two primary problems observed in samples generated via our model trained with no transformation. Namely, the square root transformation eliminated any detectable noise at the lower end of the precipitation scale, and none of its samples contained the unnatural global shift in precipitation rates. The noise elimination at the lower end of the precipitation scale is evident in Figure 4.5. Unlike in the case of no transformation, the unnatural deviation from the test distribution shape below approximately 0.2 millimetres per hour is absent in the samples generated via the square root transformation.

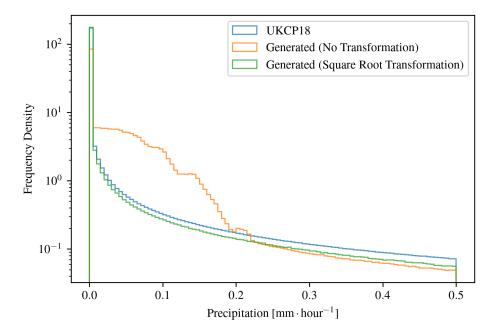


Figure 4.5: Distribution of precipitation rates below 0.5 millimetres per hour in samples generated via our model trained with the square root transformation compared to both samples generated via our model trained with no transformation and the test set from UKCP18. The format is the same as Figure 4.2

The significant improvement in sample quality is clear evidence that a transformation of the input can be critical to the performance of a diffusion model.

4.3 Learning a Transformation

4.3.1 Change of Variable Rule

Suppose we transform our observed variable \mathbf{x} via an element-wise monotonically-increasing vector-valued function $\mathbf{h}: [-1,1]^D \to [-1,1]^D$, such that:

$$\mathbf{h}(\mathbf{x}) = \begin{bmatrix} h(x_1) \\ h(x_2) \\ \vdots \\ h(x_D) \end{bmatrix}$$
(4.1)

In other words, our vector-valued function **h** applies an identical transformation $h: [-1,1] \to [-1,1]$ to each element x_i of **x**. The change of variable rule states that:

$$p_{\theta}(\mathbf{x}) = \tilde{p}_{\theta}(\mathbf{h}(\mathbf{x})) \left| \det \left(\frac{d\mathbf{h}(\mathbf{x})}{d\mathbf{x}} \right) \right|$$
 (4.2)

where det is the determinant operator, and $d\mathbf{h}(\mathbf{x})/d\mathbf{x}$ is the Jacobian of transformation \mathbf{h} . Since \mathbf{h} is an element-wise transformation, the Jacobian matrix is diagonal:

$$\frac{d\mathbf{h}(\mathbf{x})}{d\mathbf{x}} = \begin{bmatrix}
\frac{\partial \tilde{x}_1}{\partial x_1} & 0 & \cdots & 0 \\
0 & \frac{\partial \tilde{x}_2}{\partial x_2} & \cdots & 0 \\
\vdots & \vdots & \ddots & \vdots \\
0 & 0 & \cdots & \frac{\partial \tilde{x}_D}{\partial x_D}
\end{bmatrix}$$
(4.3)

The determinant of a diagonal matrix is simply the product of its diagonal elements; thus, we can write:

$$\det\left(\frac{d\mathbf{h}(\mathbf{x})}{d\mathbf{x}}\right) = \prod_{i=1}^{D} \frac{\partial h(x_i)}{\partial x_i}.$$
(4.4)

Therefore, we can rewrite the change of variables rule under transformation \mathbf{h} in the following simplified form:

$$p_{\theta}(\mathbf{x}) = \tilde{p}_{\theta}(\mathbf{h}(\mathbf{x})) \prod_{i=1}^{D} \frac{\partial h(x_i)}{\partial x_i}$$
(4.5)

We can thus formulate the negative log-likelihood of ${\bf x}$ as follows:

$$-\log p_{\theta}(\mathbf{x}) = -\log \left(\tilde{p}_{\theta}(\mathbf{h}(\mathbf{x})) \prod_{i=1}^{D} \frac{\partial h(x_i)}{\partial x_i} \right)$$
(4.6)

$$= -\log \tilde{p}_{\theta}(\mathbf{h}(\mathbf{x})) - \log \left(\prod_{i=1}^{D} \frac{\partial h(x_i)}{\partial x_i} \right)$$
(4.7)

$$= -\log \tilde{p}_{\theta}(\mathbf{h}(\mathbf{x})) - \sum_{i=1}^{D} \log \left(\frac{\partial h(x_i)}{\partial x_i} \right)$$
(4.8)

4.3.2 ELBO Loss with a Change of Variable

The ELBO loss for a transformed datapoint $\mathbf{h}(\mathbf{x})$, denoted $\tilde{\mathcal{L}}_{\mathrm{ELBO}}(\mathbf{h}(\mathbf{x}))$, serves as a variational bound on the negative log-likelihood of $\mathbf{h}(\mathbf{x})$ and is given by:

$$-\log \tilde{p}_{\theta}(\mathbf{h}(\mathbf{x})) \leq \tilde{\mathcal{L}}_{\text{ELBO}}(\mathbf{h}(\mathbf{x})) = \mathbb{E}_{\mathbf{z}_{0},\dots,\mathbf{z}_{1} \sim \tilde{q}(\mathbf{z}_{0},\dots,\mathbf{z}_{1}|\mathbf{h}(\mathbf{x}))} \left[\log \frac{\tilde{p}_{\theta}(\mathbf{z}_{0},\dots,\mathbf{z}_{1},\mathbf{h}(\mathbf{x}))}{\tilde{q}(\mathbf{z}_{0},\dots,\mathbf{z}_{1}|\mathbf{h}(\mathbf{x}))}\right]$$
(4.9)

Plugging the inequality into Equation 4.8, we obtain:

$$-\log p_{\theta}(\mathbf{x}) \le \tilde{\mathcal{L}}_{\text{ELBO}}(\mathbf{h}(\mathbf{x})) - \sum_{i=1}^{D} \log \left(\frac{\partial h(x_i)}{\partial x_i} \right)$$
(4.10)

Thus, if we were to parameterise the transformation \mathbf{h}_{ω} , with transformative parameters ω , we would theoretically be able to learn some optimal transformation to approximately minimise the negative log-likelihood of the observed variable \mathbf{x} . We denote

For completeness, the loss with transformative parameters ω yields only a minor modification to Equation 4.10 and is given by:

$$p_{\theta}(\mathbf{x}) \le \tilde{\mathcal{L}}_{\text{ELBO}}(\mathbf{h}_{\omega}(\mathbf{x})) - \sum_{i=1}^{D} \log \left(\frac{\partial h_{\omega}(x_i)}{\partial x_i} \right)$$
 (4.11)

However, as detailed in Section 2.7.8, optimising our generative parameters θ via the ELBO loss yields suboptimal results. Diffusion models in the broader literature have achieved far superior results via optimisation with the weighted loss \mathcal{L}_{WL} .

4.3.3 Weighted Loss with a Change of Variable

The λ -dependent weighting function $w(\lambda)$ in the weighted loss \mathcal{L}_{WL} causes it to lose a direct theoretical relationship with the negative log-likelihood of the observed variable \mathbf{x} . Thus, we cannot directly apply the change of variable approach to the weighted loss as we did for the ELBO loss. However, we can consider the individual contribution of the negative log-determinant term in the loss given in Equation 4.11 to derive an empirically good loss function for our diffusion model that jointly optimises the generative parameters θ and the transformative parameters ω . Importantly, since we constrain the output range of h_{ω} to be [-1,1], the neural network cannot infinitely extend the output range. As such, it has to learn some optimal monotonically increasing function to minimise:

$$-\sum_{i=1}^{D} \log \left(\frac{\partial h_{\omega}(x_i)}{\partial x_i} \right) \tag{4.12}$$

Intuitively, since the log function itself is monotonically increasing, the negative log-determinant term will encourage a steeper gradient for areas of the input space [-1,1] wherein individual elements x_i occur most often. Conversely, the term will individually encourage a shallower gradient for areas of the input space [-1,1] wherein individual elements x_i occur least often. Using this intuition, we hypothesised that we could add the negative log-determinant term to the weighted loss to jointly optimise the generative parameters θ and the transformative parameters ω to achieve higher-quality samples. We denote the weighted loss for a transformed datapoint $\mathbf{h}_{\omega}(\mathbf{x})$ by $\tilde{\mathcal{L}}_{\mathrm{WL}}(\mathbf{h}_{\omega}(\mathbf{x}))$, and thus our new loss function is given by:

$$\tilde{\mathcal{L}}_{\mathrm{WL}}(\mathbf{h}_{\omega}(\mathbf{x})) - \sum_{i=1}^{D} \log \left(\frac{\partial h_{\omega}(x_i)}{\partial x_i} \right)$$
(4.13)

While this approach arguably lacks solid theoretical justification, we found that, in practice, it yielded promising results.

4.3.4 Transformation Neural Network

We utilise a simple neural network with two fully-connected hidden layers to learn the transformation h_{ω} from Equation 4.13. The network contains only a single neuron in both the input and output layers and 16 neurons in each hidden layer; we use the tanh activation function for each neuron in both hidden layers and for the output neuron. We restrict all the weights—not including biases—to be positive to ensure the learnt transformation is monotonically increasing. Since the logarithm of zero is undefined, we add a small multiple of the input to the network's output to prevent the gradient from being zero at any point. We scale the output to the range [-1, 1].

4.3.5 Improved Weighted Loss with a Change of Variable

We introduce a masking function $m: [-1,1] \to \{0,1\}$ that takes a single element x_i of some datapoint \mathbf{x} as input and outputs a binary value, such that:

$$m(x_i) = \begin{cases} 1 & \text{if } x_i \in \mathcal{M} \\ 0 & \text{otherwise} \end{cases}$$
 (4.14)

where $\mathcal{M} \subseteq [-1,1]$ is a pre-defined masking set with no learnable parameters, which we use to assert greater control over the transformation learnt by our transformation neural network. We incorporate the masking function into our loss given in Equation 4.13 to yield:

$$\tilde{\mathcal{L}}_{WL}(\mathbf{h}_{\omega}(\mathbf{x})) - \sum_{i=1}^{D} \log \left(\frac{\partial h_{\omega}(x_i)}{\partial x_i} \right) \cdot m(x_i)$$
(4.15)

As such, we can use the masking function and masking set to influence the gradient of the learnt transformation. Only elements x_i within the masking set \mathcal{M} will contribute to the negative log-determinant term in the loss. On one end of the scale, if we were to set $\mathcal{M} = [-1, 1]$, this would equate to the loss of Equation 4.13. Conversely, if we were to set $\mathcal{M} = \emptyset$, then the log-determinant term would be redundant—it would never contribute to the loss.

Empirically, we found $\mathcal{M} = [-0.99975, 1]$ prevented the transformation network from learning too extreme a transformation. In simple terms, we prevent the elements of our training set that are extremely small—to the extent that they can be, or can almost be, considered 'dry' for all practical intents and purposes—contributing to the transformation. Importantly, we did not conduct any systematic hyperparameter search to determine some optimal masking set; thus, the masking set we used may not be optimal.

4.3.6 Comparison of the Learnt and Square Root Transformations

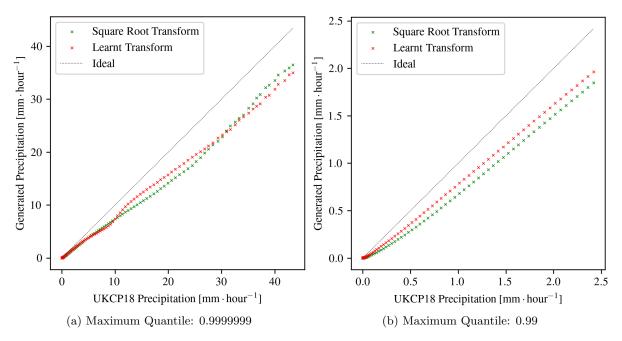


Figure 4.6: Quantile-quantile plots for individual cells within the test set from UKCP18 against both samples generated via our model trained with the square root transformation and samples generated via our model that learnt its transformation. A given point represents the precipitation rate for a given quantile in both the test set and the generated samples. Green points are for the square root transformation; red points are for the learnt transformation. The left and right plots are both for identical samples, but the left plot uses a maximum quantile of 0.9999999, while the right plot uses a maximum quantile of 0.99.

4.3.7 Samples

4.4 Temporal Interpolation

4.5 Autoregressive Generation of Longer Sequences

4.6 Forecasting

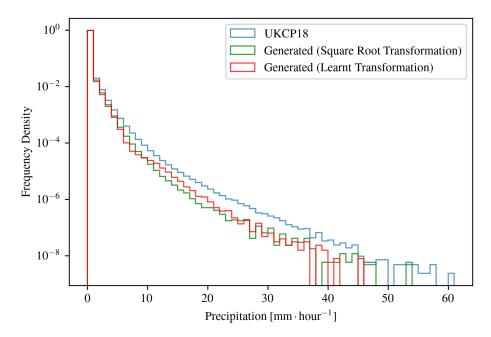
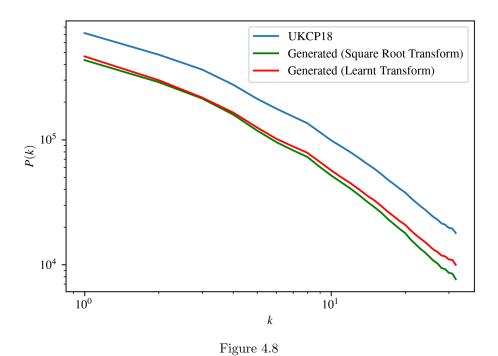
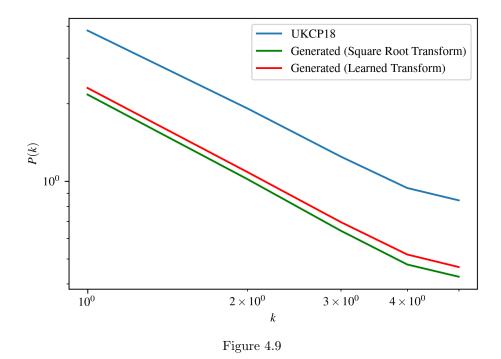


Figure 4.7: Distribution of precipitation rates in samples generated via our model trained with the square root transformation, our model that learnt its transformation, and the test set from UKCP18. The format is the same as Figure 4.3.





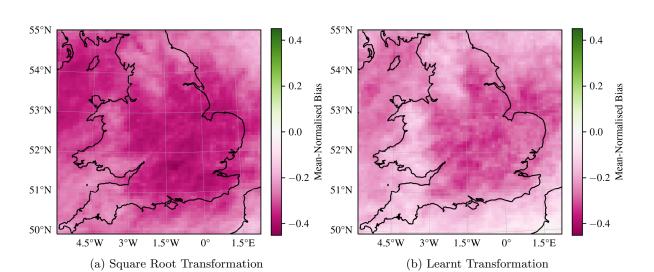


Figure 4.10: Mean-normalised bias at each cell representing the same geographical area in the samples generated via our model trained with the square root transformation (left) and our model that learnt its transformation (right). The mean absolute mean-normalised bias is 0.31 for the square root transformation and 0.23 for the learnt transformation.

Chapter 5

Conclusion

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Appendix A

Diffusion Models

A.1 Derivation of $q(\mathbf{z}_t|\mathbf{z}_s)$

From Equation 2.42, we know $q(\mathbf{z}_t|\mathbf{x})$ is an isotropic Gaussian probability density function. As such, we can sample $\mathbf{z}_t \sim q(\mathbf{z}_t|\mathbf{x})$ by sampling $\boldsymbol{\epsilon}_t \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$ from the multivariate standard Gaussian distribution and computing:

$$\mathbf{z}_t = \alpha_t \mathbf{x} + \sigma_t \boldsymbol{\epsilon}_t \tag{A.1}$$

With some algebraic manipulation, we can show that:

$$\mathbf{z}_t = \alpha_t \mathbf{x} + \sqrt{\sigma_t^2} \boldsymbol{\epsilon}_t \tag{A.2}$$

$$= \alpha_t \mathbf{x} + \sqrt{\sigma_t^2 - \frac{\alpha_t^2}{\alpha_s^2} \sigma_s^2 + \frac{\alpha_t^2}{\alpha_s^2} \sigma_s^2} \epsilon_t$$
(A.3)

$$= \alpha_t \mathbf{x} + \sqrt{\sigma_t^2 - \frac{\alpha_t^2}{\alpha_s^2} \sigma_s^2 + \left(\frac{\alpha_t}{\alpha_s} \sigma_s\right)^2} \boldsymbol{\epsilon}_t$$
 (A.4)

The sum of two independent Gaussian random variables with mean μ_1 and μ_2 and variance σ_1^2 and σ_2^2 is a Gaussian random variable with mean $\mu_1 + \mu_2$ and variance $\sigma_1^2 + \sigma_2^2$. As such, we can manipulate the above equation further to show that:

$$\mathbf{z}_t = \alpha_t \mathbf{x} + \sqrt{\sigma_t^2 - \frac{\alpha_t^2}{\alpha_s^2} \sigma_s^2} \boldsymbol{\epsilon}_t^* + \frac{\alpha_t}{\alpha_s} \sigma_s \boldsymbol{\epsilon}_s$$
 (A.5)

$$= \alpha_t \mathbf{x} + \frac{\alpha_t}{\alpha_s} \sigma_s \boldsymbol{\epsilon}_s + \sqrt{\sigma_t^2 - \frac{\alpha_t^2}{\alpha_s^2} \sigma_s^2} \boldsymbol{\epsilon}_t^*$$
(A.6)

$$= \frac{\alpha_s}{\alpha_s} \alpha_t \mathbf{x} + \frac{\alpha_t}{\alpha_s} \sigma_s \boldsymbol{\epsilon}_s + \sqrt{\sigma_t^2 - \frac{\alpha_t^2}{\alpha_s^2} \sigma_s^2} \boldsymbol{\epsilon}_t^*$$
(A.7)

$$= \frac{\alpha_t}{\alpha_s} (\alpha_s \mathbf{x} + \sigma_s \boldsymbol{\epsilon}_s) + \sqrt{\sigma_t^2 - \frac{\alpha_t^2}{\alpha_s^2} \sigma_s^2} \boldsymbol{\epsilon}_t^*$$
(A.8)

(A.9)

where ϵ_t^* , $\epsilon_s \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$ are similarly both sampled from the multivariate standard Gaussian distribution. We can substitute $\mathbf{z}_s = \alpha_s \mathbf{x} + \sigma_s \epsilon_s$ into the above equation to show that:

$$\mathbf{z}_t = \frac{\alpha_t}{\alpha_s} \mathbf{z}_s + \sqrt{\sigma_t^2 - \frac{\alpha_t^2}{\alpha_s^2} \sigma_s^2} \boldsymbol{\epsilon}_t^*$$
(A.10)

$$= \alpha_{t|s} \mathbf{z}_s + \sigma_{t|s} \boldsymbol{\epsilon}_t^* \tag{A.11}$$

$$\sim \mathcal{N}\left(\mathbf{z}_{t}; \alpha_{t|s}\mathbf{z}_{s}, \sigma_{t|s}^{2}\mathbf{I}\right)$$
 (A.12)

The subscript t|s relates to the fact that $\alpha_{t|s}$ and $\sigma_{t|s}$ define the parameters of the Gaussian probability density function $q(\mathbf{z}_t|\mathbf{z}_s)$.

A.2 α -Cosine Noise Schedule

Before truncation, the continuous-time version of the α -cosine schedule [10] as described in [6] defines α_t^2 at a given timestep $t \in [0, 1]$ as:

$$\alpha_t^2 = \cos^2\left(\frac{\pi}{2}t\right) \tag{A.13}$$

Since our model is a variance-preserving diffusion model, we can show that:

$$\sigma_t^2 = 1 - \alpha_t^2 \tag{A.14}$$

$$=1-\cos^2\left(\frac{\pi}{2}t\right)\tag{A.15}$$

$$=\sin^2\left(\frac{\pi}{2}t\right) \tag{A.16}$$

As such, we define our noise schedule before truncation \tilde{f}_{λ} for all $t \in [0,1]$ as:

$$\tilde{f}_{\lambda}(t) = \log\left(\frac{\alpha_t^2}{\sigma_t^2}\right) \tag{A.17}$$

$$= \log \left(\frac{\cos^2 \left(\frac{\pi}{2} t \right)}{\sin^2 \left(\frac{\pi}{2} t \right)} \right) \tag{A.18}$$

$$= -2\log\left(\tan\left(\frac{\pi}{2}t\right)\right) \tag{A.19}$$

However, the above noise schedule means that $\tilde{f}_{\lambda}:[0,1]\to[-\infty,\infty]$; in simpler terms, λ_t is unbounded. We follow prior work (see e.g. [6, 5]) by truncating λ_t to the desired range $[\lambda_{\min},\lambda_{\max}]$. To do so, we first need to define the inverse of the unbounded noise schedule:

$$\tilde{f}_{\lambda}^{-1}(\lambda) = \frac{2}{\pi} \arctan\left(\exp\left(-\frac{1}{2}\lambda\right)\right)$$
 (A.20)

From this, we define t_0 and t_1 as:

$$t_0 = \tilde{f}_{\lambda}^{-1}(0) = \frac{2}{\pi} \arctan\left(\exp\left(-\frac{1}{2}\lambda_{\max}\right)\right)$$
 (A.21)

$$t_1 = \tilde{f}_{\lambda}^{-1}(1) = \frac{2}{\pi} \arctan\left(\exp\left(-\frac{1}{2}\lambda_{\min}\right)\right)$$
(A.22)

The truncated noise schedule used in this work is then defined as:

$$f_{\Lambda}(t) = \tilde{f}_{\lambda}(t_0 + t(t_1 - t_0))$$
 (A.23)

$$= -2\log\left(\tan\left(\frac{\pi}{2}(t_0 + t(t_1 - t_0))\right)\right)$$
 (A.24)

A.3 v-Prediction Parameterisation

From Equation A.1, for a given datapoint $\mathbf{x} \sim q(\mathbf{x})$, we can sample latent variable $\mathbf{z}_t \sim q(\mathbf{z}_t|\mathbf{x})$ via:

$$\mathbf{z}_t = \alpha_t \mathbf{x} + \sigma_t \boldsymbol{\epsilon} \tag{A.25}$$

where $\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$ is multivariate standard Gaussian noise. We define the velocity of \mathbf{z}_t as

$$\mathbf{v}_t = \frac{d\mathbf{z}_t}{d\psi} \tag{A.26}$$

i.e. the derivative of \mathbf{z}_t with respect to ψ , which itself is:

$$\psi_t = \arctan\left(\frac{\sigma_t}{\alpha_t}\right) \tag{A.27}$$

$$= \arctan\left(\frac{\sin\left(\frac{\pi}{2}(t_0 + t(t_1 - t_0))\right)}{\cos\left(\frac{\pi}{2}(t_0 + t(t_1 - t_0))\right)}\right)$$
(A.28)

$$=\arctan\left(\tan\left(\frac{\pi}{2}(t_0+t(t_1-t_0))\right)\right) \tag{A.29}$$

$$= \frac{\pi}{2}(t_0 + t(t_1 - t_0)) \tag{A.30}$$

when using the truncated continuous-time α -cosine noise schedule as per Section 2.7.4. As such, we can formulate the velocity as:

$$\mathbf{v}_{t} = \frac{\mathbf{z}_{t}}{d\psi} = \frac{d\cos(\psi)}{d\psi}\mathbf{x} + \frac{d\sin(\psi)}{d\psi}\boldsymbol{\epsilon}$$
(A.31)

$$= -\sin(\psi)\mathbf{x} + \cos(\psi)\boldsymbol{\epsilon} \tag{A.32}$$

$$= \alpha_t \epsilon - \sigma_t \mathbf{x} \tag{A.33}$$

We can rearrange the above to derive a form for \mathbf{x} in terms of \mathbf{z}_t and \mathbf{v}_t as follows:

$$\mathbf{v}_t = -\sin(\psi)\mathbf{x} + \cos(\psi)\boldsymbol{\epsilon} \tag{A.34}$$

$$\sin(\psi)\mathbf{x} = \cos(\psi)\boldsymbol{\epsilon} - \mathbf{v}_t \tag{A.35}$$

$$= \cos(\psi) \left(\frac{\mathbf{z}_t - \cos(\psi)\mathbf{x}}{\sin(\psi)} \right) - \mathbf{v}_t$$
 (A.36)

$$\sin^{2}(\psi)\mathbf{x} = \cos(\psi)\mathbf{z}_{t} - \cos^{2}(\psi)\mathbf{x} - \sin(\psi)\mathbf{v}_{t}$$
(A.37)

$$\sin^{2}(\psi)\mathbf{x} + \cos^{2}(\psi)\mathbf{x} = \cos(\psi)\mathbf{z}_{t} - \sin(\psi)\mathbf{v}_{t}$$
(A.38)

$$(\sin^2(\psi) + \cos^2(\psi))\mathbf{x} = \cos(\psi)\mathbf{z}_t - \sin(\psi)\mathbf{v}_t$$
(A.39)

$$\mathbf{x} = \cos(\psi)\mathbf{z}_t - \sin(\psi)\mathbf{v}_t \tag{A.40}$$

$$= \alpha_t \mathbf{z}_t - \sigma_t \mathbf{v}_t \tag{A.41}$$

As per Equation A.33, during training we can

We define the velocity of \mathbf{z}_t as

We rearrange to get:

As such:

$$\mathbf{x} = \alpha_t \mathbf{z}_t - \sigma_t \mathbf{v}_t \tag{A.42}$$

During training, we train the model to minimise:

$$\mathbb{E}_{\mathbf{x},\boldsymbol{\epsilon},t} \left[\|\mathbf{v}_t - \hat{\mathbf{v}}_{\theta}(\mathbf{z}_t, \lambda_t)\|_2^2 \right] \tag{A.43}$$