Training Diffusion Models (with applications to statistics)

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Diffusion Model Recap

Perturb the data with a stochastic differential equation

$$
dX_t = \underbrace{f(t)X_t dt}_{\text{drift}} + \underbrace{g(t)dW_t}_{\text{noise}}
$$

- For SDEs of this form, the distribution of $X_t\,|\,X_0$ will be $N(m(t)X_0, s(t)^2)$ where m and s can be found by integration
- This is a generalisation of the discrete time noising processes **T** used by and [Ho et al. \[2020\]](#page-31-1) and [Song and Ermon \[2019\]](#page-33-1)

Time Reversal

■ The time reversal of the noising SDE is

$$
dX_t = [f(t)X_t - g(t)^2 \nabla_x \log p_t(X_t)] dt + g(t) dW_t
$$

- **Here, the score function** $\nabla_x \log p_t(x)$ is the gradient of the log pdf of X_t with respect to x.
- If we can approximate this, the above SDE can be used to generate new samples

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Score Matching

- **Method for approximating an unnormalised probability density** by learning its score function
- If we had access to the true score, the ideal objective would be explicit score matching:

$$
J(\theta) = \mathbb{E}_{\rho(x)}\left[\frac{1}{2} \|\psi(x;\theta) - \nabla_x \log p(x)\|_2^2\right]
$$

This minimises the MSE between the approximation $\psi(x;\theta)$ and the true score $\nabla p(x)$

Denoising Score Matching

- **Denoising score matching Vincent, 2011, is an** approximation that matches the score function of a kernel density estimate of the target density
- **Using kernel q, this can be seen as the noised data** distribution $\tilde{x} = x + e$, where $e \sim q(e)$.
- [Vincent \[2011\]](#page-33-2) showed that the following objective is equivalent to explicit score matching on the score of \tilde{x} :

$$
L_{DSM}(\theta) = \mathbb{E}_{q(x,\tilde{x})}\left[\frac{1}{2}||\psi(\tilde{x};\theta) - \nabla_{\tilde{x}}\log q(\tilde{x}-x)||_2^2\right]
$$

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Denoising Score Matching

$$
L_{DSM}(\theta) = \mathbb{E}_{p_0(x)q(\tilde{x}|x)} \left[\frac{1}{2} ||\psi(\tilde{x};\theta) - \nabla_{\tilde{x}} \log q(\tilde{x}|x) ||_2^2 \right]
$$

- **This does not require the score of the data density, only the** score of the noising kernel q.
- For example, for a Gaussian kernel $e \sim N(0,\sigma^2)$ we have $\nabla_{\tilde{\mathsf{x}}} \log q(\tilde{\mathsf{x}}|\mathsf{x}) = \frac{1}{\sigma^2}(\mathsf{x}-\tilde{\mathsf{x}})$, the direction that removes the noise from \tilde{x} .
- If we approximate p_0 with a finite sample, this matches the score of a kernel density estimate rather than the true p_t

Figure: Score matching on a warped Gaussian: target distribution, noised samples at $t = 0.15$ with direction of $\nabla \log p_{t|0}$ indicated by arrows, learned score function at $t \approx 0$.

Why use a diffusion?

- In principle, we could simply do this for some small noise level and use e.g. unadjusted Langevin dynamics to generate new samples
- There are obvious problems with this KDE works poorly for high dimensional distributions
- [Song and Ermon \[2019\]](#page-33-1) showed empirically that doing this fails to recover mode weights in mixture distributions even when the true score is used

Figure: From [Song and Ermon \[2019\]](#page-33-1): target distribution, samples using ULA with the true score function, samples using ULA with a sequence of noised densities

Time Conditioning

- **Although we are training with simple KDEs, the 'magic' of** diffusion models comes from using a sequence of noise levels, and from training a single model across time
- Score estimates are implicitly smoothed through time in a way that can give better estimates in low density regions than could be obtained with separate models
- This benefit comes from the neural network approximation choosing a good architecture is key
- So, we need an objective function that trains a single model for all noise levels
- We choose the kernel q in DSM to be the transition density $p_{t|0}(\mathsf{x}_t \, | \, \mathsf{x}_0)$ of the forward diffusion, to learn the desired ∇ log p_t
- **[Song and Ermon \[2019\]](#page-33-1) incorporate time conditioning in the** objective by taking a weighted expectation over t:

$$
\mathbb{E}_t\left\{\lambda(t)\,\mathbb{E}_{\rho_0(x)\rho_t(x_t|x)}\left[\frac{1}{2}\|\psi(x_t,t;\theta)-\nabla_{x_t}\log\rho_{t|0}(x_t\,|\,x_0)\|_2^2\right]\right\}
$$

$$
\mathbb{E}_t\left\{\lambda(t)\,\mathbb{E}_{p_0(x)p_t(x_t|x)}\left[\frac{1}{2}\|\psi(x_t,t;\theta)-\nabla_{x_t}\log p_{t|0}(x_t|x_0)\|_2^2\right]\right\}
$$

- Here, t is distributed uniformly on [0, 1], and $\lambda : \mathbb{R} \to \mathbb{R}_{>0}$ is m, a weighting function.
- [Song et al. \[2021\]](#page-33-3) showed that choosing the weighting $\lambda(t)=g(t)^2$ makes this an upper bound for the model KL divergence

Learning the noising process

- Using the SDE formulation allows us to vary the time discretisation used in sampling
- \blacksquare In continuous time, different noising processes can be equivalent to each other. This means the SDE itself does not have to be fixed in advance either
- [Kingma et al. \[2023\]](#page-32-0) reparameterise the DSM objective to show that it is invariant to changes to the VP SDE that preserve the signal-to-noise ratio at $t = 0$ and $t = 1$
- **This can be used to optimise the noise schedule to reduce the** variance of the objective function, speeding up training

(a) \log SNR vs time t

(b) Variance of VLB estimate

Figure: From [Kingma et al. \[2023\]](#page-32-0) - comparison of objective function variance for different noise schedules

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What if we know p_0 ?

- **There has been much recent interest in using diffusion models** for sampling in Bayesian statistics, even when the target density is known:
	- They are useful as surrogate models when the target density is expensive to evaluate
	- They can successfully sample from complex, multi-modal distributions, so are attractive as an alternative to MCMC

Target Score Matching

- A key difficulty is that the noised density ρ_t is typically intractable even if we know the target density p_0 , so we still need to train a score approximation
- **Denoising score matching (DSM) often struggles to** approximate the score function at low noise levels, since the variance of its score estimates explodes as $t \to 0$
- [De Bortoli et al. \[2024\]](#page-31-2) proposed an alternative objective, which uses a rescaling of the unnoised score function rather than the score of the noising distribution

Target Score Matching

The following objective can be used to estimate ∇ log p_t :

$$
L_{\text{TSM}}(\theta, t) = \mathbb{E}_{X_0, X_t} \left[\|\psi(x_t, t; \theta) - m(t)^{-1} \nabla \log p_0(x_0)\|_2^2 \right]
$$

- L_{TSM} is very well behaved near $t = 0$ where the regression target is a low variance estimator of the true score, less so for large t since typically $m(t) \rightarrow 0$
- We can get an objective that is well behaved across time by taking a weighted combination of TSM and DSM

Diffusion-Based Samplers

- **THE TSM** incorporates evaluations of the true density, but it still requires an initial sample from the target distribution to compute the objective function
- **Diffusions have desirable properties for sampling from complex** distributions (e.g. good mixing for multimodal distributions) so there has been recent interest in using them for sampling
- **For example, [Phillips et al. \[2024\]](#page-32-1) start with an initial** approximate sample, which is refined over repeated rounds of training

Motivation: Diffusion vs Tempering

Diffusion models interpolate between the target distribution and a tractable distribution, much like tempering [\[Neal, 2001\]](#page-32-2):

$$
p_t(x) = p(x)^{1-t} \phi(x)^t
$$

- Unlike diffusions, the intermediate densities ρ_t in tempering are known and do not have to be estimated
- **However, diffusions can outperform tempering on multimodal** distributions with differing mode weights

Figure: From [Phillips et al. \[2024\]](#page-32-1), comparing the intermediate densities in tempering and noising. In tempering, the mode weights can 'switch'.

Score Network

- Can in theory use any architecture that maps $\mathbb{R}^d \to \mathbb{R}^d$ with time as an input
- In practice, the network can have a huge impact on results. Try and use domain knowledge to choose an appropriate architecture, add time as an input to each layer
- For 'low dimensional' distributions ($d \approx 100$), a small feedforward MLP or ResNet will do

[Denoising Score Matching](#page-1-0) [Score Matching Variations](#page-12-0) [Neural Network Choice](#page-20-0) [References](#page-31-0)

The U-net

Figure: From [Ronneberger et al. \[2015\]](#page-32-3): image segmentation

The U-net is an architecture proposed by [Ronneberger et al. \[2015\]](#page-32-3) for image-to-image tasks, which has become ubiquitous as an architecture for score nets in image generation

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Figure: From [Ronneberger et al. \[2015\]](#page-32-3): U-net architecture

(Exact details like number/type of convolutions and inclusion of dense layers can vary) メロト マ何 トマ ミトマ ミト

- The LHS is a typical CNN architecture, with a sequence of convolutions that decrease the resolution and increase the number of channels
- The idea is for each channel to \blacksquare extract a different key feature of the image
- ■ The RHS mirrors the LHS, with transposed convolutions returning the image to its original dimensions
- This uses the extracted features to \sim construct an output image

- The low-res final output of the LHS does not contain precise information about where features are located, but we need each output pixel to relate to the corresponding input pixel
- So, the full U-net includes skip connections concatenating outputs from the LHS onto the inputs of the RHS to help with localisation

Energy based parameterisation

- **Since the score function of a distribution determines its** density up to normalising constant, we can use score matching to estimate the target density directly
- This idea was proposed by [Salimans and Ho \[2021\]](#page-32-4) as a way of ensuring that the score approximation is in fact a valid score function
- **This is known as an energy-based model (EBM)** because we model an energy function $E(x, t; \theta)$ and approximate p_t by $\exp(-E(x, t; \theta))$

A common parameterisation is:

$$
E(x, t; \theta) = \frac{1}{2s(t)} ||x - \psi(x, t; \theta)||_2^2,
$$

where $\psi(x,t;\rho):\mathbb{R}^d\rightarrow\mathbb{R}^d$ is a neural network and $s(t)^2$ is the variance of the noising kernel $p_{t|0}$

- **■** The gradient $-\nabla_x E(x,t;\rho)$ is substituted into the usual score matching objective in training
- **[Salimans and Ho \[2021\]](#page-32-4) found that this performed similarly** but no better than the usual parameterisation

Composing diffusion models

[Du et al. \[2023\]](#page-31-3) found that using an EBM to perform MCMC sampling enables sampling from compositions of diffusion models

Figure: From [Du et al. \[2023\]](#page-31-3): sampling from product and mixture distributions, where a reverse SDE is not available

[Denoising Score Matching](#page-1-0) [Score Matching Variations](#page-12-0) [Neural Network Choice](#page-20-0) [References](#page-31-0)

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Figure: [Du et al. \[2023\]](#page-31-3) - classifier guidance as sampling from a product distribution

Figure: [Du et al. \[2023\]](#page-31-3) - using product distributi[ons](#page-28-0) t[o](#page-30-0) [co](#page-28-0)[mb](#page-29-0)[i](#page-30-0)[ne](#page-19-0)[pro](#page-33-0)[m](#page-19-0)[p](#page-20-0)[ts](#page-33-0)

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Applications in Statistics

- **E** Sampling from products of posterior distributions can be used in Bayesian statistics to sample from the posterior conditioned on their pooled datasets
- Simulation-based inference [\[Geffner et al., 2023\]](#page-31-4) approximating single-observation posteriors requires fewer simulator calls than conditioning jointly on larger datasets
- Divide-and-conquer MCMC [\[Trojan et al., 2024\]](#page-33-4) if the full dataset is very large, it can be computationally intractable to sample directly from the full posterior distribution

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