

Diffusion Probabilistic Models

- Concurrently, a similar to score-matching class of models: diffusion models
- Diffusion models also define a forward and reverse diffusion process, where $t = 0$ corresponds to the data distribution, and $t = T$ a unit-Gaussian distribution

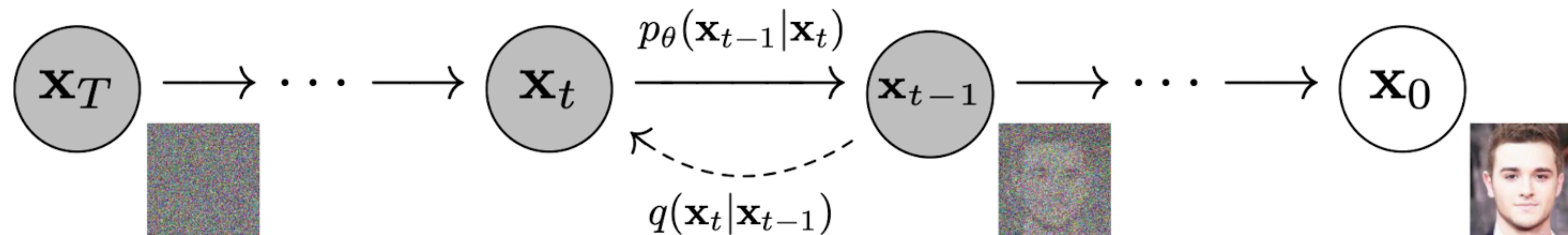


Figure 2: The directed graphical model considered in this work.

Diffusion probabilistic models, Sohl-Dickstein et al., 2015

Denosing diffusion probabilistic models, Ho et al., 2020

Diffusion models beat GANs on image synthesis, 2021

<https://lilianweng.github.io/posts/2021-07-11-diffusion-models/>

Forward diffusion process

- In forward diffusion we add small Gaussian noise to our data till it looks like isotropic Gaussian

$$q(\mathbf{x}_t | \mathbf{x}_{t-1}) = \mathcal{N}(\mathbf{x}_t; \sqrt{1 - \beta_t} \mathbf{x}_{t-1}, \beta_t \mathbf{I}), \quad q(\mathbf{x}_{1:T} | \mathbf{x}_0) = \prod_{t=1}^T q(\mathbf{x}_t | \mathbf{x}_{t-1})$$

- We can define the conditional distribution at any time step t w.r.t. step $t = 0$

$$\begin{aligned} \mathbf{x}_t &= \sqrt{a_t} \mathbf{x}_{t-1} + \sqrt{1 - a_t} \mathbf{z}_{t-1} && , \text{ where } \mathbf{z}_{t-1}, \mathbf{z}_{t-2}, \dots \sim \mathcal{N}(0, 1) \\ &= \sqrt{a_t a_{t-1}} \mathbf{x}_{t-2} + \sqrt{1 - a_t a_{t-1}} \bar{\mathbf{z}}_{t-2} && , \text{ where } \bar{\mathbf{z}}_{t-2} \text{ merges two Gaussians} \\ &= \dots \\ &= \sqrt{\bar{a}_t} \mathbf{x}_0 + \sqrt{1 - \bar{a}_t} \bar{\mathbf{z}} \end{aligned}$$

$$q(\mathbf{x}_t | \mathbf{x}_0) = \mathcal{N}(\mathbf{x}_t; \sqrt{\bar{a}_t} \mathbf{x}_0, \sqrt{1 - \bar{a}_t} \mathbf{I})$$

- For this we use that when merging two Gaussians, we get another Gaussian with variance $\sigma_1^2 + \sigma_2^2$

Reverse diffusion process

- Efficiently parameterising reverse diffusion we can combine with variational inference

$$L_{VLB} = L_T + L_{T-1} + \dots + L_0$$

where $L_T = D_{KL}(q(\mathbf{x}_T | \mathbf{x}_0) || p_\theta(\mathbf{x}_T))$

$$L_t = D_{KL}(q(\mathbf{x}_t | \mathbf{x}_{t+1}, \mathbf{x}_0) || p_\theta(\mathbf{x}_t | \mathbf{x}_{t+1})) \text{ for } 1 \leq t \leq T - 1$$

$$L_0 = -\log p_\theta(\mathbf{x}_0 | \mathbf{x}_1)$$

- Since we have Gaussian distributions the KL terms can be computed in closed form
- L_T does not depend any parameters and it can be dropped
- L_0 depends on the final decoder output

Parameterising L_t

- By smart parameterisation of the Gaussians, learning boils down to minimising

$$L_t^{\text{simple}} = \mathbb{E}_{\mathbf{x}_0, \epsilon_t} \left[\left\| \epsilon_t - \epsilon_\theta \left(\sqrt{\bar{a}_t} \mathbf{x}_0 + \sqrt{1 - \bar{a}_t} \epsilon_t, t \right) \right\|_2^2 \right]$$

- The model **learns to predict** the **noise** added to the **transformed signal!!!!**

Algorithm 1 Training

- 1: **repeat**
- 2: $\mathbf{x}_0 \sim q(\mathbf{x}_0)$
- 3: $t \sim \text{Uniform}(\{1, \dots, T\})$
- 4: $\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$
- 5: Take gradient descent step on
 $\nabla_\theta \left\| \epsilon - \epsilon_\theta(\sqrt{\bar{\alpha}_t} \mathbf{x}_0 + \sqrt{1 - \bar{\alpha}_t} \epsilon, t) \right\|_2^2$
- 6: **until** converged

Algorithm 2 Sampling

- 1: $\mathbf{x}_T \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$
 - 2: **for** $t = T, \dots, 1$ **do**
 - 3: $\mathbf{z} \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$ if $t > 1$, else $\mathbf{z} = \mathbf{0}$
 - 4: $\mathbf{x}_{t-1} = \frac{1}{\sqrt{\alpha_t}} \left(\mathbf{x}_t - \frac{1 - \alpha_t}{\sqrt{1 - \bar{\alpha}_t}} \epsilon_\theta(\mathbf{x}_t, t) \right) + \sigma_t \mathbf{z}$
 - 5: **end for**
 - 6: **return** \mathbf{x}_0
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Example trajectories

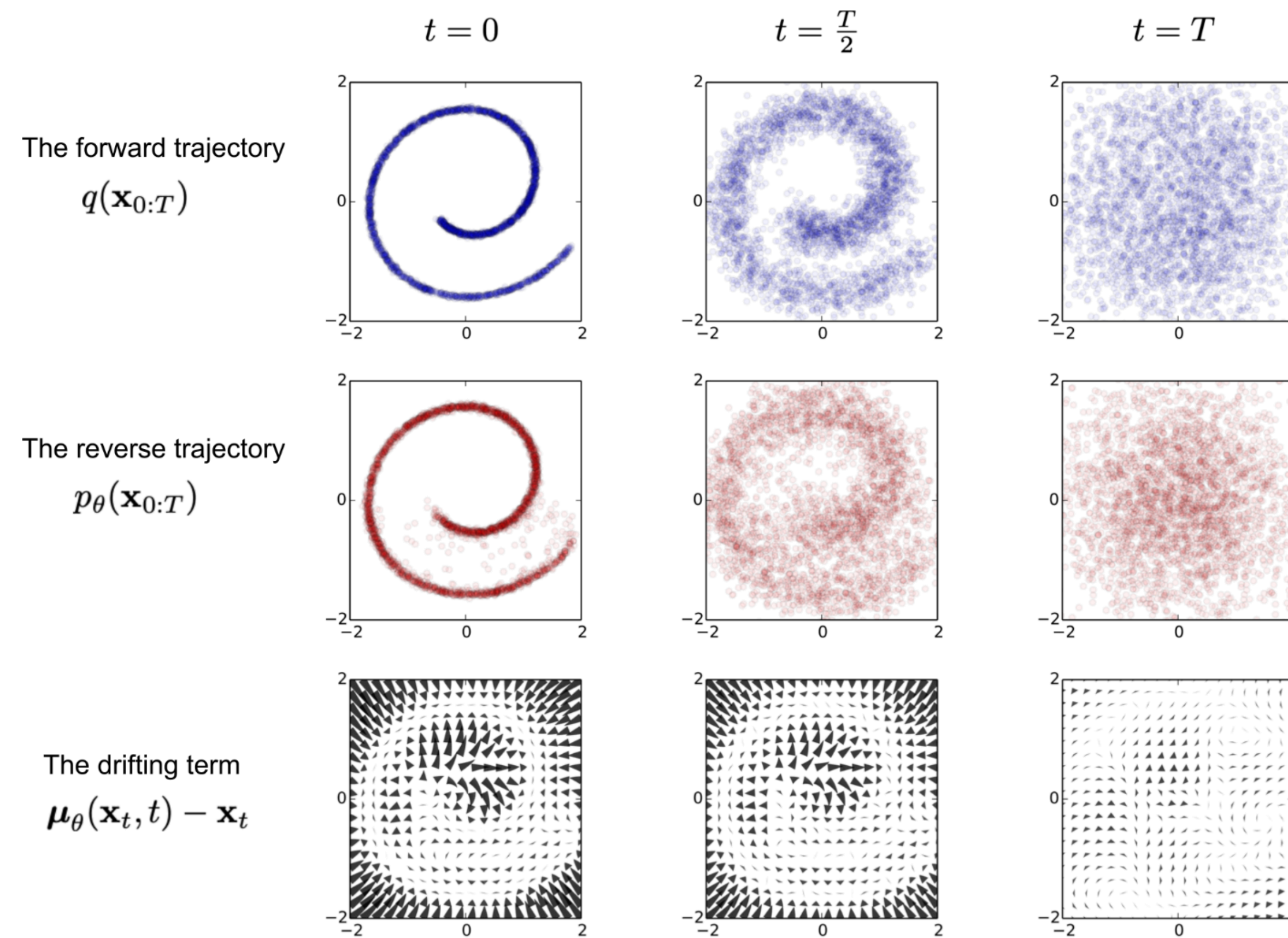


Fig. 3. An example of training a diffusion model for modeling a 2D swiss roll data. (Image source: [Sohl-Dickstein et al., 2015](#))

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Score-matching & Diffusion Generative Models

Qualitative results

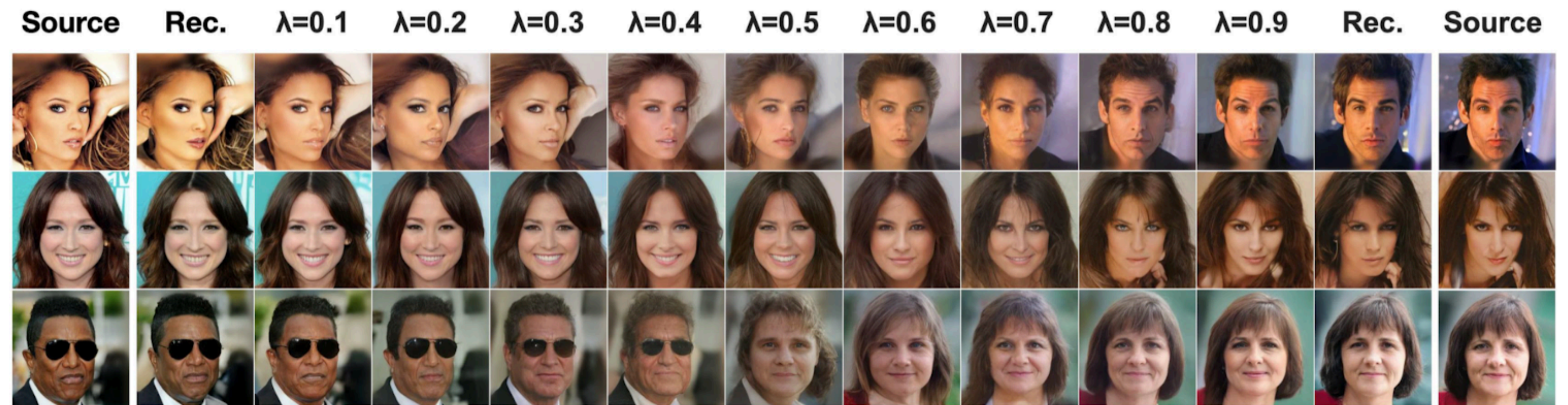
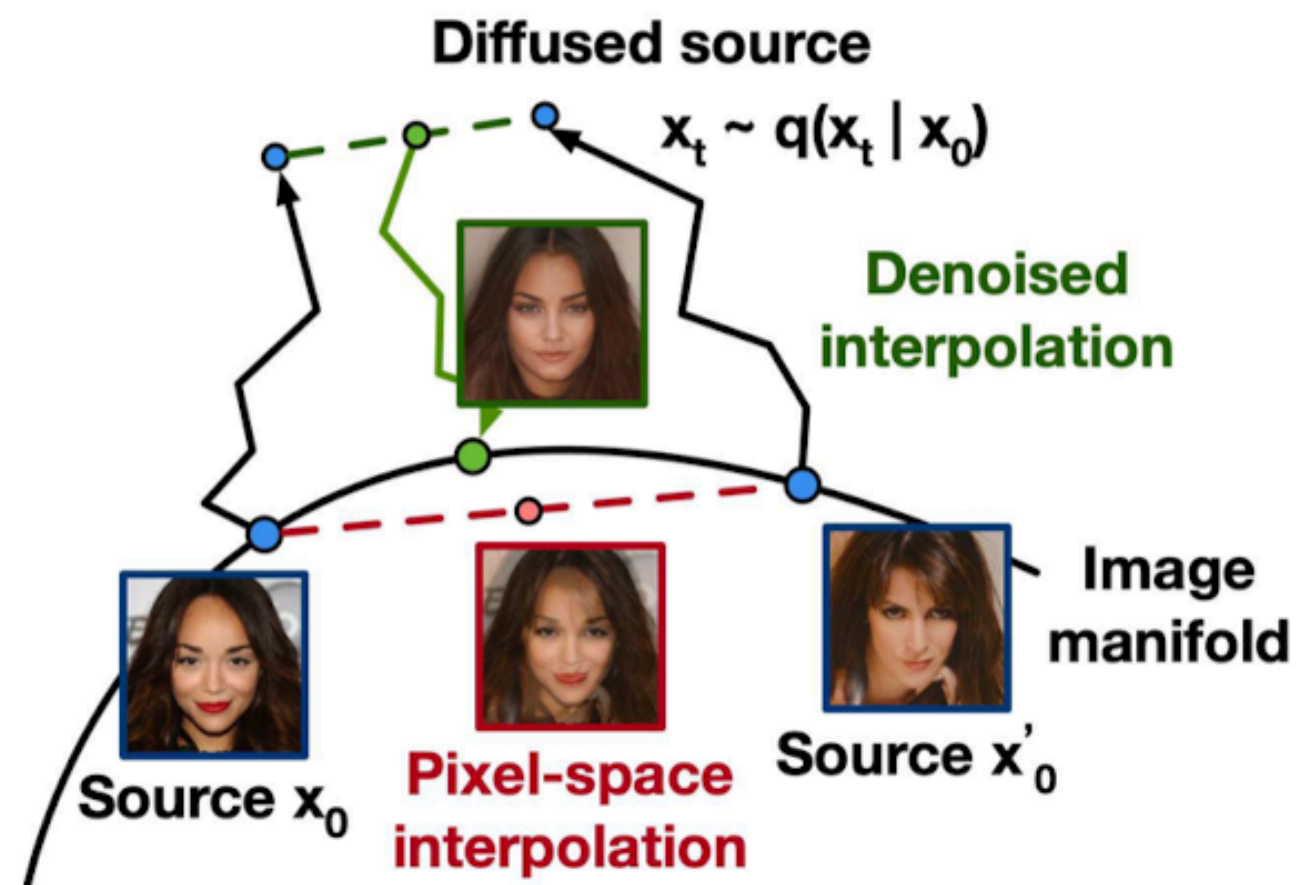


Figure 8: Interpolations of CelebA-HQ 256x256 images with 500 timesteps of diffusion.

Denoising diffusion probabilistic models, Ho et al., 2020

Take-home message

- Diffusion/score-matching models are both tractable and flexible
- However, they are still quite slow to sample from compared to GANs
- The reason is that they require very long chains of time steps up to $T = 1,000$
- Great opportunities for learning the data structure effectively and efficiently enough
- Promising results in modelling inverse problems