### 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

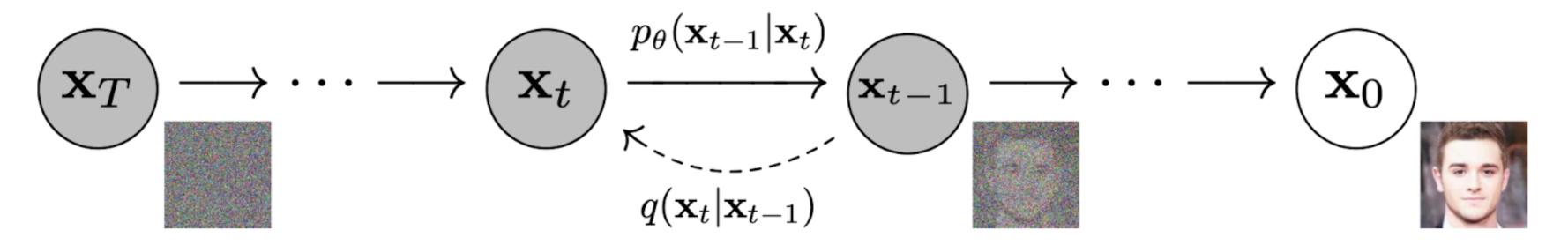


Figure 2: The directed graphical model considered in this work.

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

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

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

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

• 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

$$\begin{split} L_{VLB} &= L_T + L_{T-1} + \ldots + L_0 \\ \text{where } L_T &= D_{KL} \big( q(\mathbf{x}_T | \mathbf{x}_0) || p_{\theta}(\mathbf{x}_T) \big) \\ L_t &= D_{KL} \big( q(\mathbf{x}_t | \mathbf{x}_{t+1}, \mathbf{x}_0) || p_{\theta}(\mathbf{x}_t | \mathbf{x}_{t+1}) \big) \text{ for } 1 \leq t \leq T-1 \\ L_0 &= -\log p_{\theta}(\mathbf{x}_0 | \mathbf{x}_1) \end{split}$$

- 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

$$L_{t}^{\text{simple}} = \mathbb{E}_{\mathbf{x}_{0}, \boldsymbol{\epsilon}_{t}} \left[ \left\| \boldsymbol{\epsilon}_{t} - \boldsymbol{\epsilon}_{\theta} \left( \sqrt{\bar{a}_{t}} \mathbf{x}_{0} + \sqrt{1 - \bar{a}_{t}} \boldsymbol{\epsilon}_{t}, t \right) \right\|_{2}^{2} \right]$$

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

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### 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

$$\nabla_{\theta} \left\| \boldsymbol{\epsilon} - \boldsymbol{\epsilon}_{\theta} (\sqrt{\bar{\alpha}_{t}} \mathbf{x}_{0} + \sqrt{1 - \bar{\alpha}_{t}} \boldsymbol{\epsilon}, t) \right\|^{2}$$

6: **until** converged

#### **Algorithm 2** Sampling

- 1:  $\mathbf{x}_T \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$
- 2: **for** t = T, ..., 1 **do**
- 3:  $\mathbf{z} \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$  if t > 1, else  $\mathbf{z} = \mathbf{0}$
- 4:  $\mathbf{x}_{t-1} = \frac{1}{\sqrt{\alpha_t}} \left( \mathbf{x}_t \frac{1-\alpha_t}{\sqrt{1-\bar{\alpha}_t}} \boldsymbol{\epsilon}_{\theta}(\mathbf{x}_t, t) \right) + \sigma_t \mathbf{z}$
- 5: end for
- 6: return  $x_0$

### Example trajectories

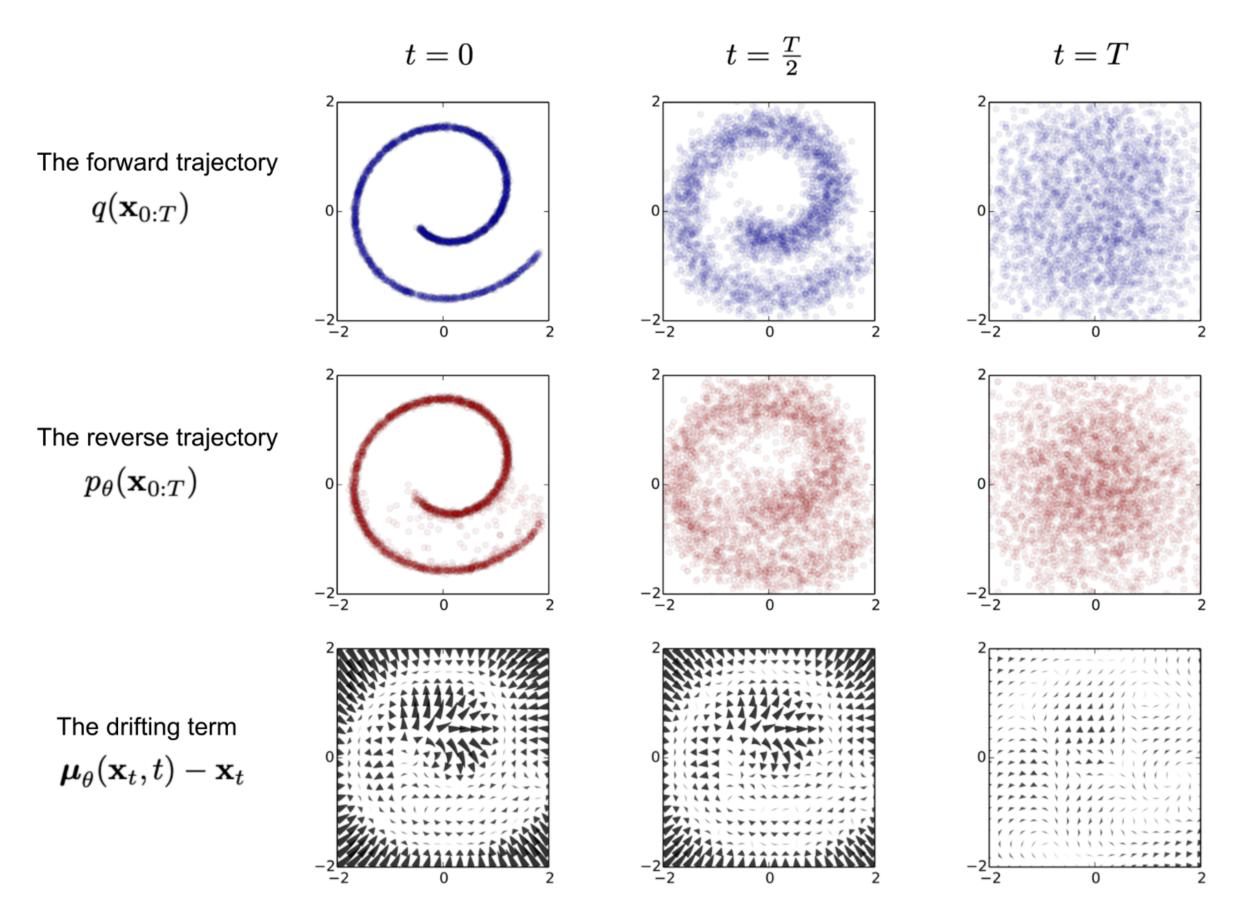


Fig. 3. An example of training a diffusion model for modeling a 2D swiss roll data. (Image source: Sohl-Dickstein et al., 2015)

### Qualitative results

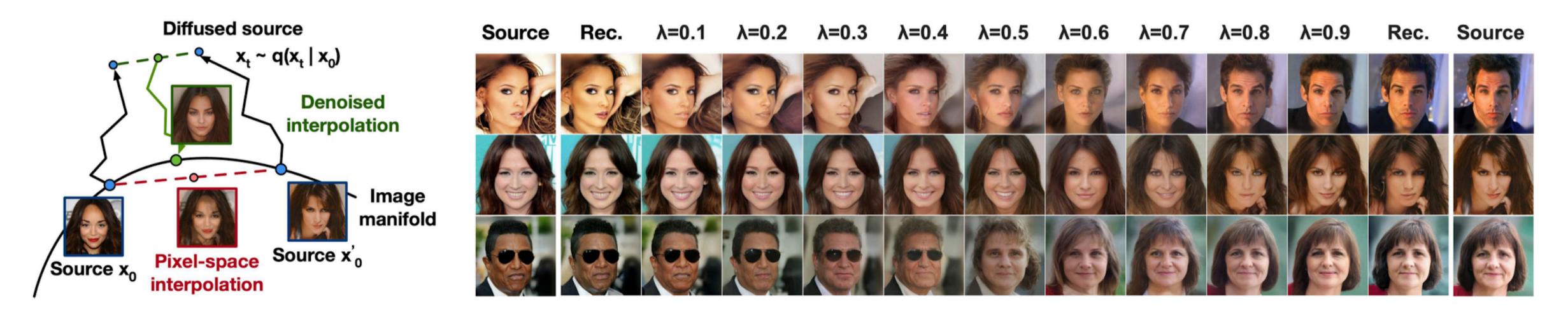


Figure 8: Interpolations of CelebA-HQ 256x256 images with 500 timesteps of diffusion.

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