Deep Unsupervised Learning using Nonequilibrium Thermodynamics

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0.1. Background

- Probabilistic models suffer from a tradeoff between two conflicting objectives: tractability and flexibility.
- Tractable models: can be analytically evaluated and easily fit to data(e.g. a Gaussian or Laplace), but unable to aptly describe structure in rich datasets.
- Flexible models: can be molded to fit structure in arbitrary data, but evaluating, training, or drawing samples from such flexible models typically requires a very expensive Monte Carlo process.
- Inspired by non-equilibrium statistical physics, we develop an approach that simultaneously achieves both flexibility and tractability.
 - ▶ 1.Systematically and slowly destroy structure in a data distribution through an iterative forward diffusion process.
 - 2.Learn a reverse diffusion process that restores structure in data, yielding a highly flexible and tractable generative model of the data.

0.2. Intro

Method:

- We build a generative Markov chain which converts a simple known distribution (e.g. a Gaussian) into a target (data) distribution using a diffusion process. (explicitly define the probabilistic model as the endpoint of the Markov chain.)
- Since each step in the diffusion chain has an analytically evaluable probability, the full chain can also be analytically evaluated.
- Estimating small perturbations is more tractable than explicitly describing the full distribution with a single, non-analytically-normalizable, potential function.
- Since a diffusion process exists for any smooth target distribution, this method can capture data distributions of arbitrary form

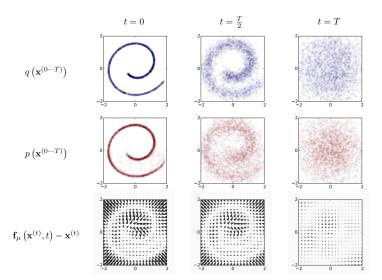
Goal:

- Define a forward diffusion process which converts any complex data distribution into a simple, tractable distribution
- Learn a finite-time reversal of this diffusion process which defines our generative model distribution

Advantages:

- extreme flexibility in model structure
- exact sampling
- (easy multiplication with other distributions, e.g. in order to compute a posterior)
- the model log likelihood, and the probability of individual states, to be cheaply evaluated

e.g. Swiss Roll



1.1 Process

1. Forward trajectory

The data distribution $q\left(x^{(0)}\right)$ is gradually converted into a well-behaved (analytically tractable) distribution $\pi(y)$ by repeated application of a Markov diffusion kernel $T_{\pi}(y|y';\beta)$ for $\pi(y)$, where β is the diffusion rate,

$$\pi(y) = \int dy' T_{\pi}(y|y';\beta)\pi(y') \tag{1}$$

$$q\left(x^{(t)}|x^{(t-1)}\right) = T_{\pi}\left(x^{(t)}|x^{(t-1)};\beta_{t}\right)$$
(2)

$$q(x^{(0\cdots T)}) = q(x^{(0)}) \prod_{t=1}^{T} q(x^{(t)}|x^{(t-1)})$$
(3)

2. Reverse trajectory

The generative distribution will be trained to describe the same trajectory, but in reverse,

$$p\left(x^{(T)}\right) = \pi\left(x^{(T)}\right) \tag{4}$$

$$p\left(x^{(0\cdots T)}\right) = p\left(x^{(T)}\right) \prod_{t=1}^{T} p\left(x^{(t-1)}|x^{(t)}\right). \tag{5}$$

1.2. Model Probability

The probability the generative model assigns to the data is

$$\label{eq:posterior} \textit{p}\left(\mathbf{x}^{(0)}\right) = \int \textit{d}\mathbf{x}^{(1\cdots\textit{T})}\textit{p}\left(\mathbf{x}^{(0\cdots\textit{T})}\right)$$

But the integral is intractable.

Inspired by **annealed importance sampling** and the **Jarzynski equality**, we instead compute:

$$p\left(x^{(0)}\right) = \int dx^{(1\cdots T)} \frac{p\left(x^{(0\cdots T)}\right)}{q\left(x^{(1\cdots T)} \mid x^{(0)}\right)} q\left(x^{(1\cdots T)} \mid x^{(0)}\right)$$

$$= \int dx^{(1\cdots T)} q\left(x^{(1\cdots T)} \mid x^{(0)}\right) \frac{p\left(x^{(0\cdots T)}\right)}{q\left(x^{(1\cdots T)} \mid x^{(0)}\right)}$$

$$= \int dx^{(1\cdots T)} q\left(x^{(1\cdots T)} \mid x^{(0)}\right) \cdot \frac{p\left(x^{(T)}\right) \prod_{t=1}^{T} p\left(x^{(t-1)} \mid x^{(t)}\right)}{\prod_{t=1}^{T} q\left(x^{(t)} \mid x^{(t-1)}\right)}.$$
(6)

This can be evaluated rapidly by averaging over samples from the forward trajectory $q\left(x^{(1\cdots T)}\mid x^{(0)}\right)$.

1.3. Training

We want to maximize $p(x_0)$ when $x_0 \sim q(x_0)$, i.e. to maximize model log likelihood (equivalently, minimize the cross entropy) $E_{x_0 \sim q(x_0)}[log(p(x_0))]$

$$L = \int dx^{(0)} q(x^{(0)}) \log p(x^{(0)})$$

$$= \int dx^{(0)} q(x^{(0)}) \cdot \log \left[\int dx^{(1\cdots T)} q(x^{(1\cdots T)} | x^{(0)}) \cdot \frac{p(x^{(T)}) \prod_{t=1}^{T} p(x^{(t-1)} | x^{(t)})}{\prod_{t=1}^{T} q(x^{(t)} | x^{(t-1)})} \right]$$

$$\geq \int dx^{(0\cdots T)} q(x^{(0\cdots T)}) \cdot \log \left[\frac{p(x^{(T)}) \prod_{t=1}^{T} p(x^{(t-1)} | x^{(t)})}{q(x^{(t)} | x^{(t-1)})} \right]$$

This reduces to $L \ge K$ (ELBO in variational inference)

$$\begin{split} K &= -\sum_{t=2}^{T} \int dx^{(0)} dx^{(t)} q\left(x^{(0)}, x^{(t)}\right) \cdot D_{\mathsf{KL}}\left(q\left(x^{(t-1)} | x^{(t)}, x^{(0)}\right) \parallel p\left(x^{(t-1)} | x^{(t)}\right)\right) \\ &+ H_{q}\left(X^{(T)} | X^{(0)}\right) - H_{q}\left(X^{(1)} | X^{(0)}\right) - H_{p}\left(X^{(T)}\right). \end{split}$$

Where the entropies and KL divergences can be analytically computed.



1.3. Training (cont.)

► Training consists of finding the reverse Markov transitions which maximize this lower bound on the log likelihood

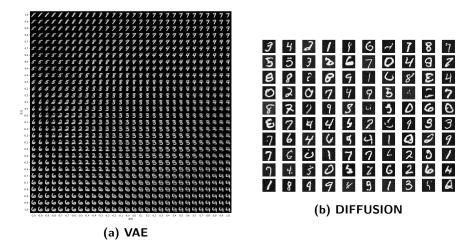
$$\hat{p}\left(x^{(t-1)} \mid x^{(t)}\right) = \underset{p(x^{(t-1)} \mid x^{(t)})}{\operatorname{argmax}} K$$

Thus, the task of estimating a probability distribution has been reduced to the task of performing regression on the functions which set the mean and covariance of a sequence of Gaussians

- ▶ Remaning: Setting The Diffusion Rate β_t
 - ▶ Gaussian diffusion: learn the forward diffusion $\beta_{2...T}$ by gradient ascent on K. The variance β_1 of the first step is fixed to a small constant to prevent overfitting. The dependence of samples from $q\left(x^{(1\cdots T)}|x^{(0)}\right)$ on $\beta_{1...T}$ is made explicit by using frozen noise as in the noise is treated as an additional auxiliary variable, and held constant while computing partial derivatives of K with respect to the parameters.
 - ▶ Binomial diffusion: Discrete state space, gradient ascent with frozen noise impossible. Instead choose the forward diffusion schedule $\beta_{1...T}$ to erase a constant fraction $\frac{1}{T}$ of the original signal per diffusion step, yielding a diffusion rate of $\beta_t = (T-t+1)^{-1}$.
 - ▶ Recent experiments suggest that it is just as effective for Gaussian diffusion to instead use the same fixed t schedule as for binomial diffusion

analytically tractable distribution
$$\pi\left(x^{(T)}\right) = \begin{cases} \mathcal{N}\left(x^{(T)}; \mathbf{0}, \mathbf{I}\right) & (\mathsf{Gauss}) \\ \mathcal{B}\left(x^{(T)}; 0.5\right) & (\mathsf{Bin}) \end{cases}$$
 Forward diffusion kernel
$$q\left(x^{(t)} \mid x^{(t-1)}\right) = \begin{cases} \mathcal{N}\left(x^{(t)}; x^{(t-1)}\sqrt{1-\beta_t}, \mathbf{I}\beta_t\right) & (\mathsf{Gauss}) \\ \mathcal{B}\left(x^{(t)}; x^{(t-1)}(1-\beta_t) + 0.5\beta_t\right) & (\mathsf{Bin}) \end{cases}$$
 Reverse diffusion kernel
$$p\left(x^{(t-1)} \mid x^{(t)}\right) = \begin{cases} \mathcal{N}\left(x^{(t-1)}; f_{\mu}\left(x^{(t)}, t\right), f_{\Sigma}\left(x^{(t)}, t\right)\right) & (\mathsf{Gauss}) \\ \mathcal{B}\left(x^{(t-1)}; f_{b}\left(x^{(t)}, t\right)\right), f_{\Sigma}\left(x^{(t)}, t\right) & (\mathsf{Bin}) \end{cases}$$
 Training targets
$$\begin{cases} f_{\mu}\left(x^{(t)}, t\right), f_{\Sigma}\left(x^{(t)}, t\right), \beta_{1\cdots T} & (\mathsf{Gauss}) \\ f_{b}\left(x^{(t)}, t\right), \beta_{1\cdots T} & (\mathsf{Bin}) \end{cases}$$
 Forward distribution
$$q\left(x^{(0\cdots T)}\right) = q\left(x^{(0)}\right) \prod_{t=1}^{T} q\left(x^{(t)} \mid x^{(t-1)}\right)$$
 Reverse distribution
$$p\left(x^{(0\cdots T)}\right) = \pi\left(x^{(T)}\right) \prod_{t=1}^{T} p\left(x^{(t-1)} \mid x^{(t)}\right)$$
 Log likelihood
$$\mathcal{L} = \int \mathrm{d}x^{(0)} q\left(x^{(0)}\right) \log p\left(x^{(0)}\right)$$
 Lower bound on log likelihood
$$\mathcal{L} = \int \mathrm{d}x^{(0)} q\left(x^{(0)}\right) - H_{q}\left(x^{(t-1)} \mid x^{(t)}, x^{(0)}\right) + P_{q}\left(x^{(t-1)} \mid x^{(t)}\right)$$

e.g. MNIST



unlike many MNIST sample figures, diffusion-generated are true samples rather than the mean of the Gaussian or binomial distribution from which samples would be drawn.

Thank you for your listening.