Linear and deep latent variable models from PPCA to VAEs

Pierre-Alexandre Mattei





Recap on directed graphical models

What's a graph, mathematically?

Definition (directed graph). A graph is a pair G = (V, E) comprising a set V of vertices or nodes together with a set $E \subset V \times V$ of edges or arcs.

$$V_{1} \longrightarrow V_{2} \longrightarrow V_{3} \qquad V = \{v_{1}, v_{2}, v_{3}, v_{4}\}$$

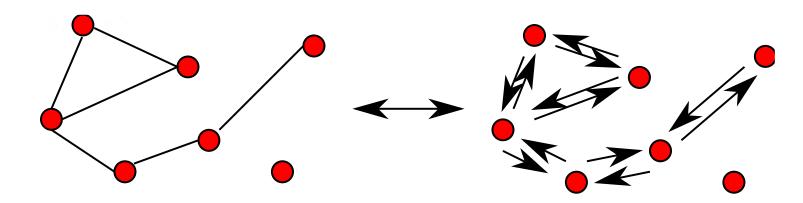
$$E = \{(v_{1}, v_{2}), (v_{2}, v_{3}), (v_{2}, v_{4}), (v_{3}, v_{4})\}$$

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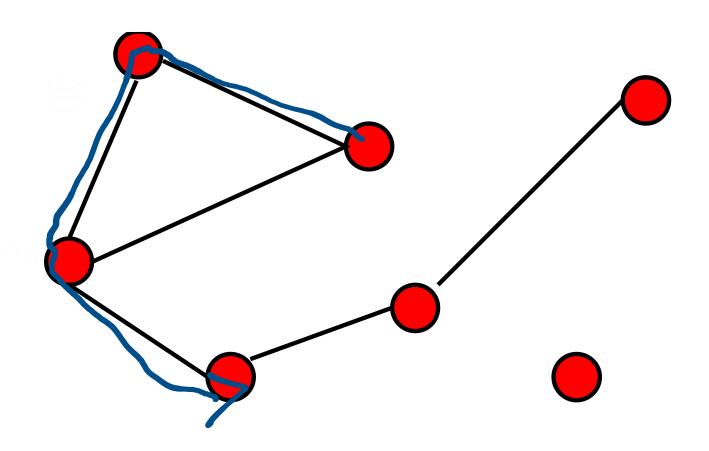
Definition (undirected graph). G = (V, E) is **undirected** if all edges go in both directions: for all $(u, v) \in V \times V$ such that $u \neq v$, we have:

$$(u,v) \in E \iff (v,u) \in E.$$



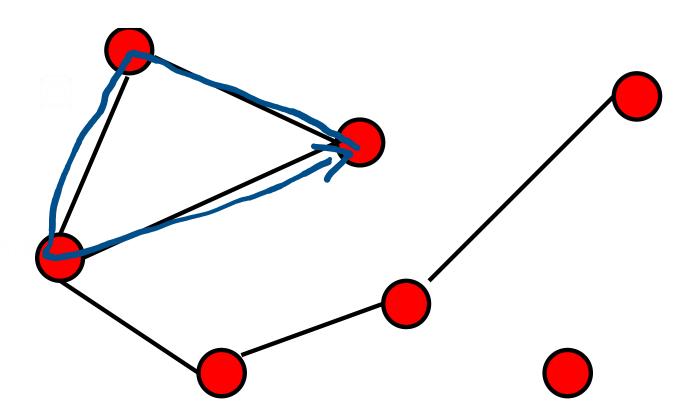
What's a path?

Definition. A path is a sequence of connected vertices that are globally distinct.



What's a cycle?

Definition. A path that begins and end at the same point is called a **cycle**.

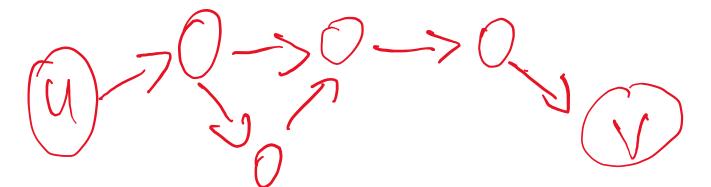


Some definitions specific to directed graphs

Definition. u is a **parent** of v if $(u, v) \in E$. We also say that v is a **children** of u.

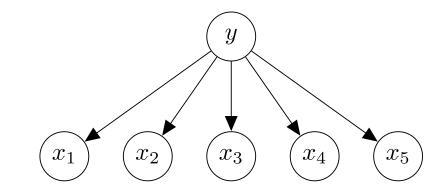


Definition. u is an **ancestor** of v if there exists a path from u to v. We also say that v is a **descendant** of u.

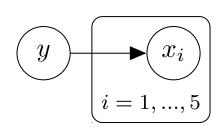


The useful plate notation

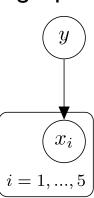
Often we deal with graphs with recurring stucture, for instance



• The plate notation is a more compact way of drawing the same graph:



or



A key concept: DAGs

Definition (DAG). A directed acyclic graph (DAG) is a directed graph without any cycle.

What are directed graphical models?

- Let G = (V, E) be a DAG whose nodes are denoted $V = \{1, ..., d\}$
- Let $X = (X_1, ..., X_d)$ be a random variable with density $p(x) = p(x_1, ..., x_d)$.
- Every node of the graph corresponds to a random variable (one vertex for each feature)

Definition (Directed graphical model). We say that p factorises in G (denoted $p \in \mathcal{L}(G)$) when, for all x,

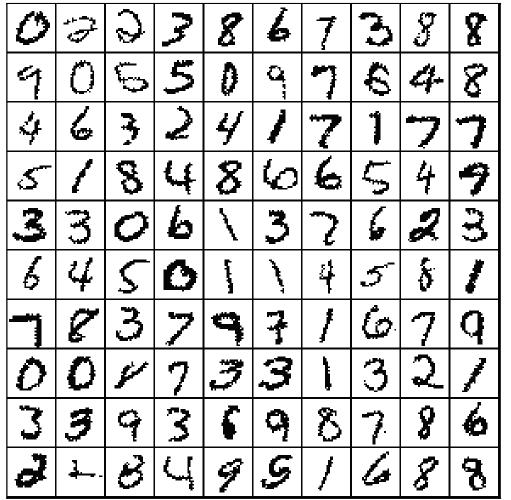
$$p(x) = \prod_{i=1}^{d} p(x_i | x_{\text{pa}_i}),$$

where, for all node i, pa_i denotes the set of parents of node i.

More examples of directed models and notations

Recurring « fil rouge » example

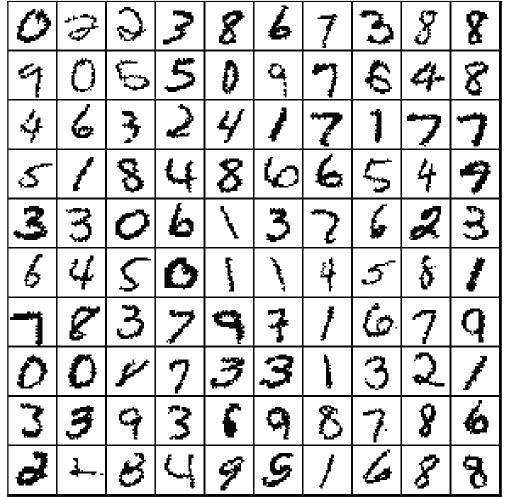
- We will often consider as a working example a binary version of MNIST
- $x_1, ..., x_n \in \mathcal{X}$ $\mathcal{X} = \{0, 1\}^{28 \times 28}$
- $y_1, ..., y_n \in \{0, ..., 9\}$



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- Binarised by Hugo Larochelle
- Sometimes called « statical binary MNIST » in VAE papers



A basic example of parametrised distribution for binary MNIST

- Let's go back to our binary images, that live in $\mathcal{X} = \{0,1\}^{28 \times 28}$
- A basic model would be to assume that each pixel corresponds to a different coin flip!

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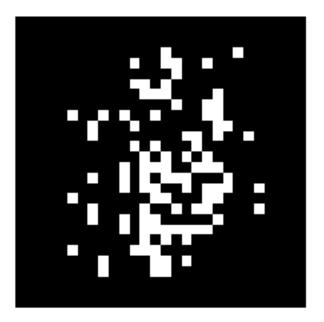
This may be rewritten

$$p_{\theta}(x_1, \dots, x_n) = \prod_{i=1}^{n} p_{\theta}(x_i) = \prod_{i=1}^{n} \prod_{j=1}^{28 \times 28} \mathcal{B}(x_{ij} | \theta_j)$$

Why is this model not great?

The MLE is just
$$\hat{\theta}_j = \frac{1}{n} \sum_{i=1}^n x_{ij}$$

Here is a sample from the model...



The factorisation is

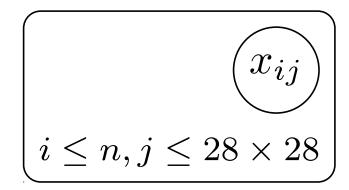
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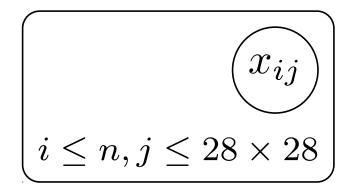
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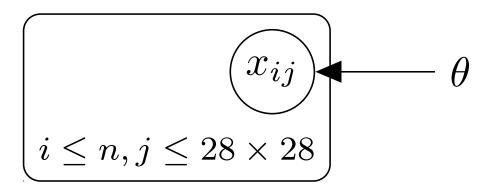


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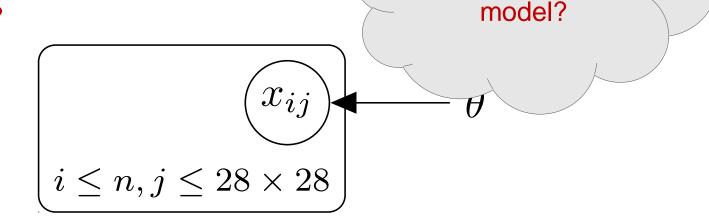


...aka "trivial graph" (because it has no edges)

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 How do we improve this weak

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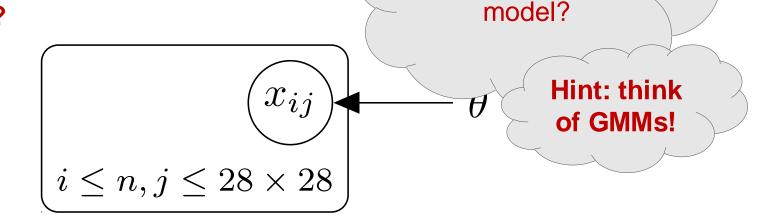


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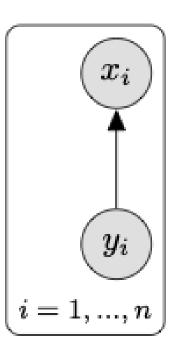
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Adding a node that represents the class improves the model!

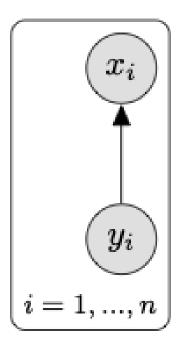
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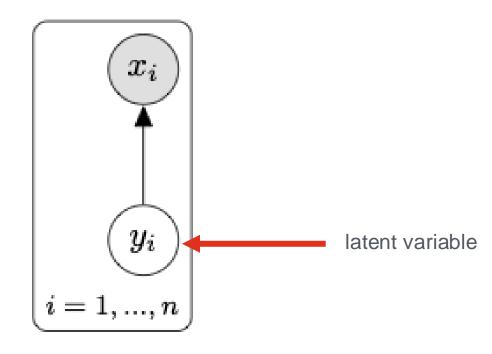


But this is a supervised model... How to make it unsupervised?

$$\theta = (\pi, (\mu_{yj})_{y \in \{0, \dots, 9\}, j \le 784})$$

Latent class analysis (aka mixtures of products of Bernoullis)

$$p_{\theta}(x_1,\ldots,x_n,y_1,\ldots,y_n) = \prod_{i=1}^n p_{\theta}(y_i) p_{\theta}(x_i|y_i) \text{ with } p_{\theta}(y) = \mathcal{M}(y|1,\pi) \text{ and } p_{\theta}(x|y) = \mathcal{B}(x|\mu_{yj})$$



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The likelihood of the supervised model is just a sum of log-densities of simple distributions:

$$\log p_{ heta}(x_1,\ldots,x_n,y_1,\ldots,y_n) = \sum_{i=1}^n \left(\log p_{ heta}(y_i) + \log p_{ heta}(x_i|y_i)
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What would be ways to maximise it? EM, gradient descent, etc.

3

Technical aparté 1

Sampling with graphical models: ancestral sampling

Why sampling from a graphical model

1. The samples are useful by themselves

Images generated by a GAN



Brock, Donahue, and Simonyan (ICLR 2019)

Why sampling from a graphical model

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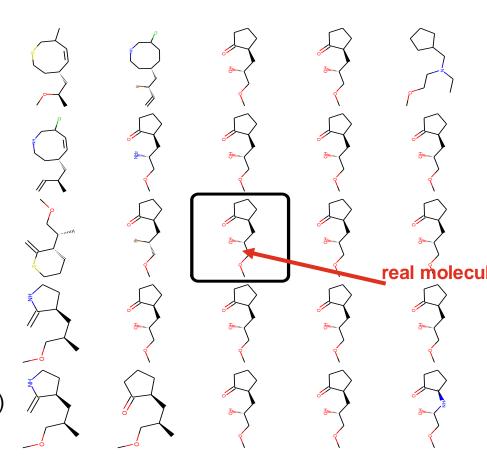
Images generated by a VAE Vahdat and Kautz (NeurlPS 2020)

Designing new molecules with a VAE

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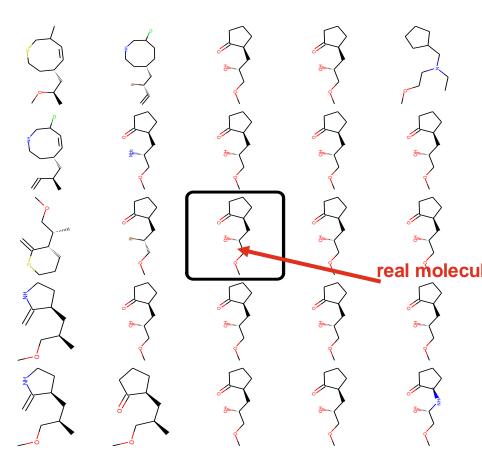


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$$\int f(x)p(x)dx \approx \frac{1}{n} \sum_{i=1}^{n} f(x_i)$$

Monte Carlo basics

- Until now, we have written integrals as sums that we can compute exhaustively, but this is
 quite rarely the case.
- When this integral is an expected value over something we know how to sample from, we can use simple Monte Carlo

Algorithm 1 Simple Monte Carlo

1: Draw $X^{(1)}, ..., X^{(n)} \overset{i.i.d.}{\sim} p$

2: $\hat{\mu} = \frac{1}{n} \sum_{i=1}^{n} f(X^{(i)})$

• Why are we expecting this to work?

Monte Carlo basics

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2:
$$\hat{\mu} = \frac{1}{n} \sum_{i=1}^{n} f(X^{(i)})$$

Proposition (Law of Large Numbers).

$$\hat{\mu} \xrightarrow{a.s.} \mu$$
 iff μ is finite

Proposition (Central Limit Theorem). If $Var(f(X)) = \sigma^2 < \infty$, then

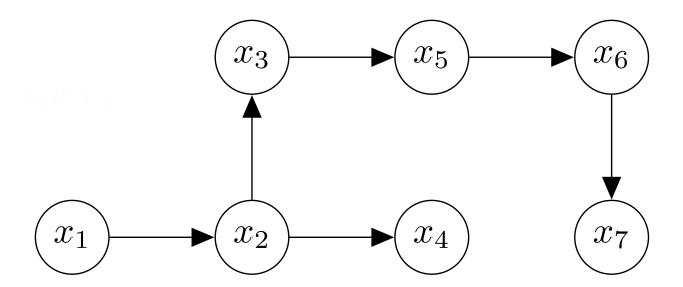
$$\sqrt{n}(\hat{\mu} - \mu) \xrightarrow{\mathcal{D}} \mathcal{N}(0, \sigma^2)$$

Algorithm 1 Ancestral sampling with topological ordering

```
for i=1 to d do

Draw z_i from p(X_i=.|X_{\mathrm{pa}_i}=z_{\mathrm{pa}_i})
end for

return (z_1,\ldots,z_d)
```



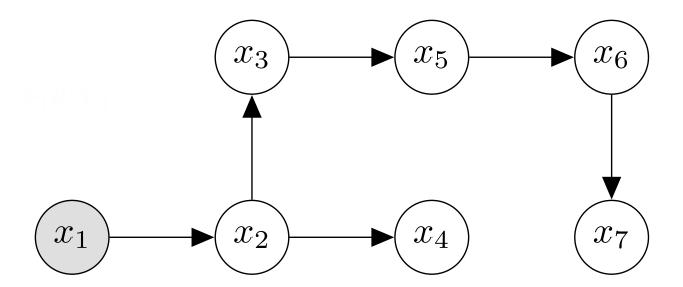
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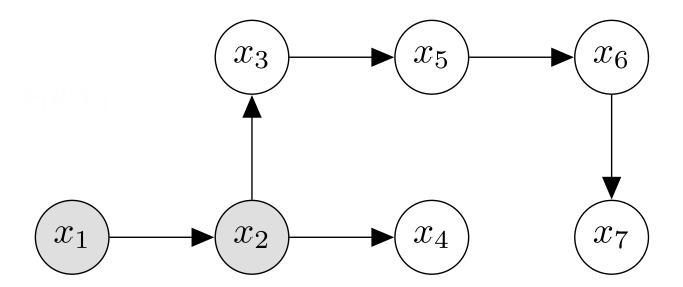
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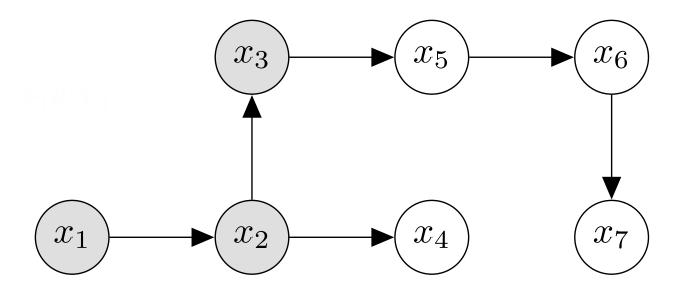
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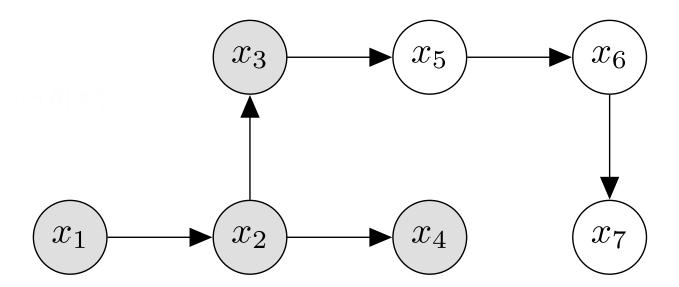


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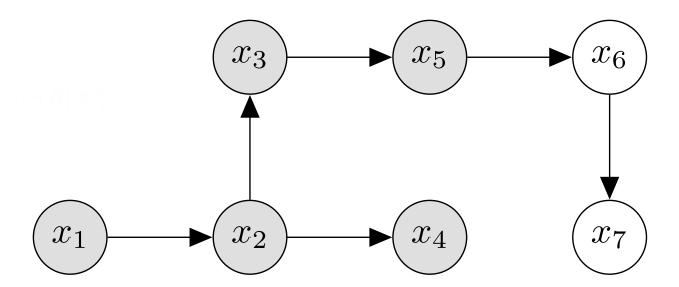
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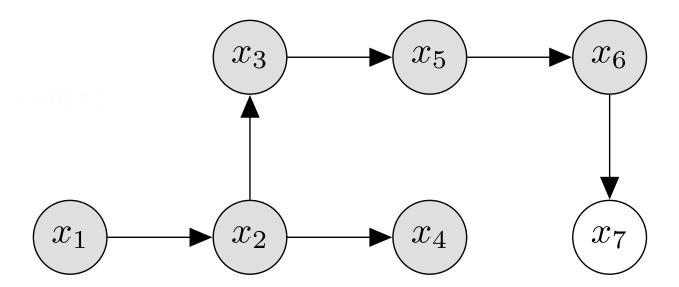
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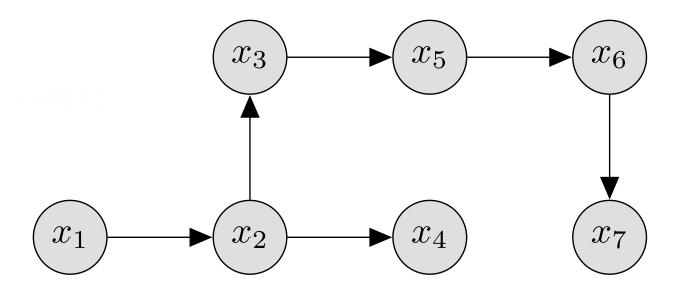
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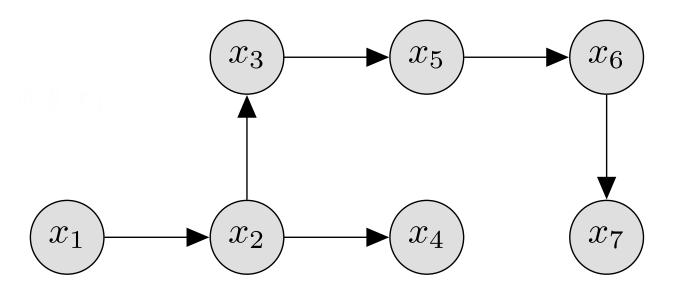
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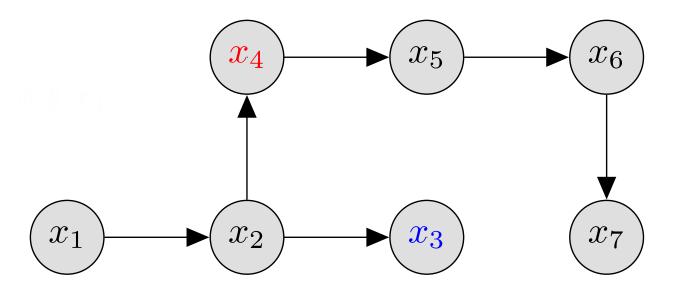
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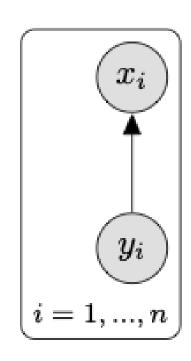
Algorithm 1 Ancestral sampling with topological ordering for i = 1 to d do Draw z_i from $p(X_i = .|X_{pa_i} = z_{pa_i})$ end for

Proposition (ancestral sampling). Algorithm 1 provides samples from the joint distribution $p(x_1, ..., x_d)$.

Proof. By induction on the topological ordering.

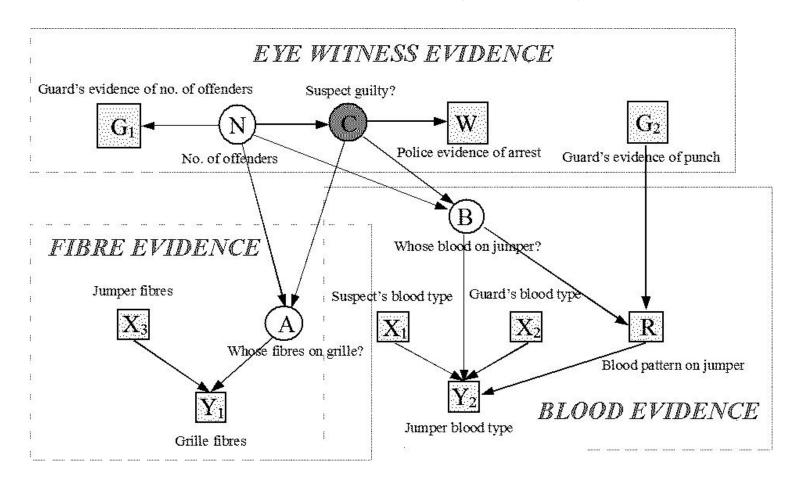
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Think of ways to implement this for some of the graphs we saw



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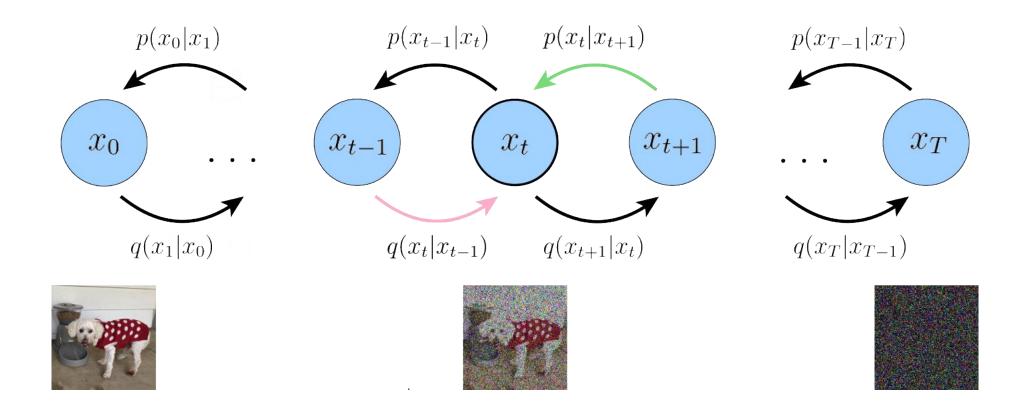
• The big graph from the forensic science example... Here the topological ordering is related to the time of the events...



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Think of ways to implement this for some of the graphs we saw

Diffusion model (again, two DAGs!)



Linear latent variable models
PPCA and friends

Why latent-variable models?

Let's assume that we have i.i.d. data $x_1, ..., x_n$ that live in a **high-dimensional space** \mathcal{X} (for example images of newspaper articles).

Often, it is reasonable to assume that these data essentially depend on a few important factors of variation:

- if we have images of faces: size of nose, color of hair, glasses or not...
- if we have newspaper articles: topics of the paper, style...
- if we have molecules: physical properties, geometrical shape...

This assumption is the cornerstone of **latent variable models**, that assume that the data are governed by **unobserved random variables** $\mathbf{z}_1, ..., \mathbf{z}_n$ **that live in a low dimensional space** \mathcal{Z} (for example \mathbf{R}^2). We can think of \mathbf{z}_i as a **code** summarizing the essential factors of the data point \mathbf{x}_i .

Linear latent-variable models

In this slide, we assume that the data are continuous (i.e. $\mathbf{x}_1,...,\mathbf{x}_n \in \mathbb{R}^p$).

Factor analysis is probably one of the oldest latent variable models (studied since at least the 1940s). The generative process is:

- $\mathbf{z}_i \sim \mathcal{N}(0, \mathbf{I}_d)$,
- $\mathbf{x}_i | \mathbf{z}_i \sim \mathcal{N}(\mathbf{W}\mathbf{z}_i + \boldsymbol{\mu}, \boldsymbol{\Psi}).$

Probabilistic PCA (PPCA, Tipping and Bishop, JRSSB, 1999) is a a slightly less general model

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 - > For such models, performing maximum likelihood is not too hard:
 - EM algorithm for FA
 - Closed-form MLE for PPCA (based on the SVD of the data matrix)

From linear (« shallow ») to deep variable models

How do we transform this kind of model

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...into "something deep"?

 $\mathbb{P}(\theta|X)$

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...into "something deep"?

We can replace the affine function $\mathbf{z} \mapsto \mathbf{W}\mathbf{z} + \boldsymbol{\mu}$ by a neural net!

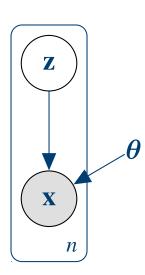
That's the key idea of deep latent variable models (DLVMs), present in particular in both variational autoencoders (VAEs), invented independently by Kingma and Welling (ICLR 2014) and Rezende, Mohamed & Wierstra (ICML 2014), and generative adversarial networks (GANs) (Goodfellow et al., NeurIPS 2014).

Deep latent variable models

Deep latent variable models

Assume that $(\mathbf{x}_i, \mathbf{z}_i)_{i \le n}$ are i.i.d. random variables driven by the model:

$$\begin{cases} \mathbf{z} \sim p(\mathbf{z}) & \text{(prior)} \\ \mathbf{x} \sim p_{\theta}(\mathbf{x} \mid \mathbf{z}) = \Phi(\mathbf{x} \mid f_{\theta}(\mathbf{z})) & \text{(observation model)} \end{cases}$$



where

- $\mathbf{z} \in \mathbb{R}^d$ is the **latent** variable,
- $x \in \mathcal{X}$ is the **observed** variable.
 - the function $f_{\theta}: \mathbb{R}^d \to H$ is a (deep) neural network called the decoder
 - $(\Phi(\cdot \mid \boldsymbol{\eta}))_{\boldsymbol{\eta} \in H}$ is a parametric family called the **observation model**, usually **very simple**: unimodal and fully factorised (e.g. multivariate Gaussians or products of multinomials)

Deep latent variable models: the role of the prior

As in regular factor analysis, the prior distribution of the latent variable is often an isotropic Gaussian $p(\mathbf{z}) = \mathcal{N}(\mathbf{z}|\mathbf{0}_d, \mathbf{I}_d)$.

Note that this prior is not a prior in the Bayesian sense (i.e., about parameter uncertainty).

Deep latent variable models: the role of the observation model

The observation model $(\Phi(\cdot \mid \boldsymbol{\eta}))_{\boldsymbol{\eta} \in H}$ usually **very simple**: unimodal and fully factorised (e.g. multivariate Gaussians or products of multinomials)

Its parameters are the output of the decoder.

$$\begin{cases} \mathbf{z} \sim p(\mathbf{z}) & \text{(prior)} \\ \mathbf{x} \sim p_{\theta}(\mathbf{x} \mid \mathbf{z}) = \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_{\theta}(\mathbf{z}), \boldsymbol{\Sigma}_{\theta}(\mathbf{z})) & \text{(Gaussian observation model)} \end{cases}$$

$$\begin{cases} \mathbf{z} \sim p(\mathbf{z}) & \text{(prior)} \\ \mathbf{x} \sim p_{\theta}(\mathbf{x} \mid \mathbf{z}) = \mathcal{B}(\mathbf{x} \mid \boldsymbol{\pi}_{\theta}(\mathbf{z})) & \text{(Bernoulli observation model)} \end{cases}$$

$$\begin{cases} \mathbf{z} \sim p(\mathbf{z}) & \text{(prior)} \\ \mathbf{x} \sim p_{\theta}(\mathbf{x} \mid \mathbf{z}) = \mathsf{St}(\mathbf{x} \mid \boldsymbol{\mu}_{\theta}(\mathbf{z}), \boldsymbol{\Sigma}_{\theta}(\mathbf{z}), \boldsymbol{\nu}_{\theta}(\mathbf{z})) & \text{(Student's t observation model)} \end{cases}$$

Deep latent variable models: back to PPCA

$$p(z) = \mathcal{N}(z|0,I_Q)$$
 $p_{ heta}(x|z) = \mathcal{N}(x|Wz + \mu,\sigma^2 I_D)$

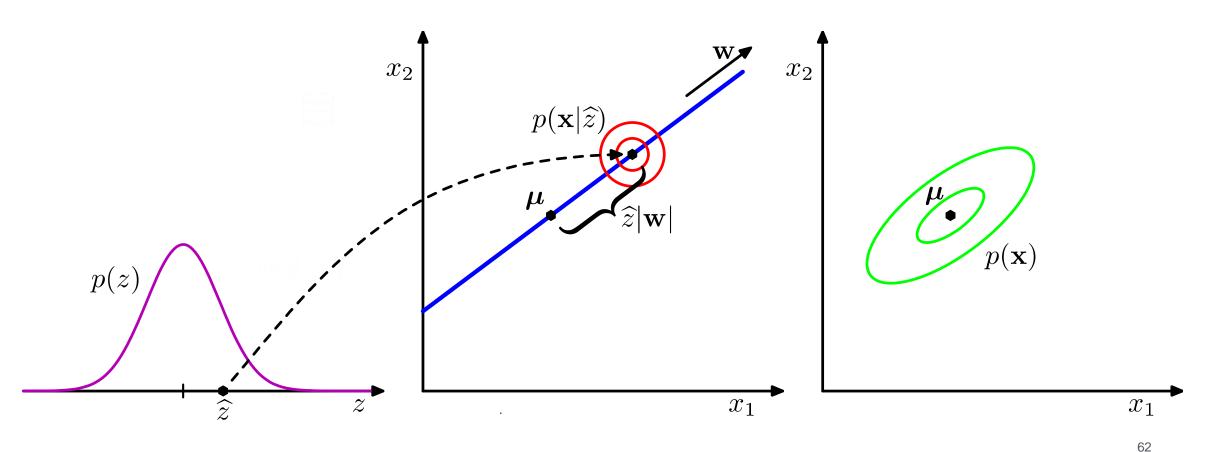


Fig. 12.9 in Bishop's Pattern Recognition and Machine Learning

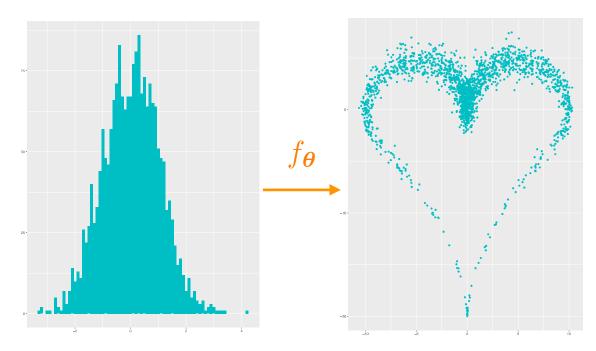
Deep latent variable models: the role of the decoder

The role of the **decoder** $f_{\theta}: \mathbb{R}^d \to H$ is:

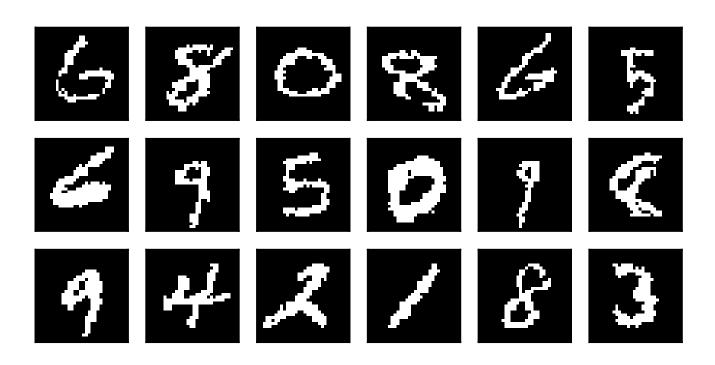
- to transform \mathbf{z} (the code) into parameters $\eta = f_{\theta}(\mathbf{z})$ of the observation model $\Phi(\cdot \mid \eta)$.
- The weights θ of the **decoder** are learned.

Simple non-linear decoder (d = 1, p = 2): $f_{\theta}(z) = \mu_{\theta}(\mathbf{z}), \Sigma_{\theta}(\mathbf{z})$ with, for all $z \in \mathbb{R}$,

$$\boldsymbol{\mu}_{\boldsymbol{\theta}}(z) = \left(10\sin(z)^3, 10\cos(z) - 10\cos(z)^4\right), \ \boldsymbol{\Sigma}_{\boldsymbol{\theta}}(\mathbf{z}) = \mathsf{Diag}\left(\left(\frac{\sin(z)}{3z}\right)^2, \left(\frac{\sin(z)}{z}\right)^2\right).$$



Training data $\{x_1, \ldots, x_n\}$ binary MNIST



Generative model for $\mathbf{z} \in \mathbb{R}^2$ and $\mathbf{x} \in \{0, 1\}^{28 \times 28}$

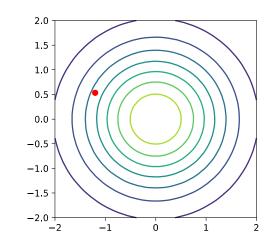
$$\begin{cases} \mathbf{z} \sim \mathcal{N}\left(\begin{bmatrix} 0 \\ 0 \end{bmatrix}, \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}\right) \\ x^{j,k} \sim \text{Bernoulli}(p = f^{j,k}(\mathbf{z})) \end{cases}$$

Decoder network

$$f(\mathbf{z}) = \text{Sigmoid}(\mathbf{V} \tanh(\mathbf{W}\mathbf{z} + \mathbf{b}) + \boldsymbol{\beta})$$

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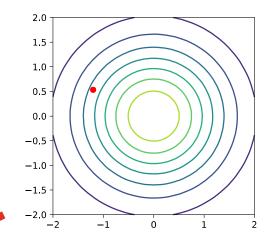
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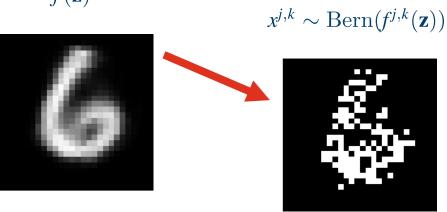
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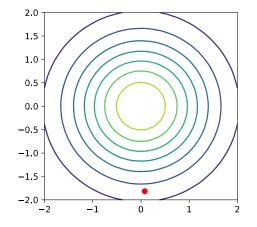
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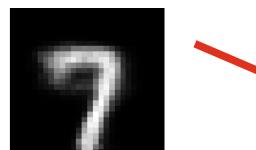
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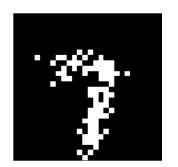
$$\mathbf{z} \sim \mathcal{N}\left(\begin{bmatrix} 0 \\ 0 \end{bmatrix}, \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} \right)$$

$$\mathbf{z} = (0.0791, -1.8165)$$





$$x^{j,k} \sim \operatorname{Bern}(f^{j,k}(z))$$



Maximum likelihood for DLVM

Given a data matrix $\mathbf{X} = (\mathbf{x}_1, \dots, \mathbf{x}_n)^\intercal \in \mathcal{X}^n$, the **log-likelihood function** for a DLVM is

$$\ell(\boldsymbol{\theta}) = \log p_{\boldsymbol{\theta}}(\mathbf{X}) = \sum_{i=1}^{n} \log p_{\boldsymbol{\theta}}(\mathbf{x}_i),$$

where

$$p_{m{ heta}}(\mathbf{x}_i) = \int_{\mathbb{R}^d} p_{m{ heta}}(\mathbf{x}_i \mid \mathbf{z}) p(\mathbf{z})$$

We would like to find a MLE $\hat{\theta} \in \operatorname{argmax}_{\theta} \ell(\theta)$.

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Maximum likelihood for DLVMs via importance sampling

A general and scalable framework to tackle these issues was proposed by Kingma & Welling (2014), Rezende et al. (2014), leading to the **variational autoencoder (VAE)**.

Here, I am going to derive this approach in a slightly different manner, largely inspired by the following paper:



Burda, Grosse & Salakhutdinov (2016), Importance weighted autoencoders, ICLR 2016

The main idea is to use Monte Carlo techniques to approximate the intractable integrals

$$p_{\boldsymbol{\theta}}(\mathbf{x}_i) = \int_{\mathbb{R}^d} p_{\boldsymbol{\theta}}(\mathbf{x}_i \mid \mathbf{z}) p(\mathbf{z}) d\mathbf{z}.$$

Aparté: Beyond simple MC: importance sampling

We have to use Monte Carlo! Our goal is to approximate an integral

$$I = \int_{\Omega} f(x)p(x)dx$$

We already saw the simple MC estimate

$$I \approx \frac{1}{K} \sum_{k=1}^{K} f(x_k) = \hat{I}_K.$$

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- This estimate has nice properties:
 - Unbiasedness:
 - ightharpoonup Consistency: $\mathbb{E}[\hat{I}_K] = I$
 - ightharpoonup Asymptotic no \hat{r} mal \hat{r} iyI

- One way of assessing the accuracy of any unbiased estimate is by looking at its variance.
 The lower the variance of an unbiased estimate, the better.
- If the samples are iid, then the variance of simple MC will be $\mathbb{V}[\hat{I}_K] = rac{1}{K}\mathbb{V}[f(x_1)]$

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- If the samples are iid, then the variance of simple MC will be $\mathbb{V}[\hat{I}_K] = rac{1}{K}\mathbb{V}[f(x_1)]$
- So the variance gets smaller and smaller at speed 1/K, that's good news!
- But it can still be pretty big, depending on the value of $V[f(x_1)]$

Can we reduce the variance of the MC estimate?

• Key idea: rather than sampling from p(x), we're going to sample from another density q(x) that we'll call a **proposal**

$$x_1,...,x_K \sim q$$

 But the integral is an expected value with respect to p. Can we turn an expected value with respect to p into an expected value with respect to q?

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$$x_1,...,x_K \sim q$$

 But the integral is an expected value with respect to p. Can we turn an expected value with respect to p into an expected value with respect to q? Yes!

$$I = \int_{\Omega} f(x)p(x)dx$$

$$= \int_{\Omega} \frac{f(x)p(x)}{q(x)} q(x)dx \approx \frac{1}{K} \sum_{k=1}^{K} \frac{f(x_k)p(x_k)}{q(x_k)} = \hat{I}_K^q.$$

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\dots and I is precisely the thing we want to compute!

- In practice, we won't be able to find this optimal proposal, but this shows that the improvements
 of importance sampling can be potentially huge!
- This simple result is therefore a motivation for looking for good proposals.

We want to approximate

$$p_{m{ heta}}(\mathbf{x}_i) = \int_{\mathbb{R}^d} p_{m{ heta}}(\mathbf{x}_i \mid \mathbf{z}) p(\mathbf{z}) \mathbf{z}$$

Idea: use importance sampling! Let $\mathbf{z}_{i1},...,\mathbf{z}_{iK}$ follow some proposal q_i :

$$\int_{\mathbb{R}^d} p_{\boldsymbol{\theta}}(\mathbf{x}_i \mid \mathbf{z}) p(\mathbf{z}_{ik}) d\mathbf{z} \approx \frac{1}{K} \sum_{k=1}^K \frac{p_{\boldsymbol{\theta}}(\mathbf{x}_i \mid \mathbf{z}_{ik}) p(\mathbf{z})}{q_i(\mathbf{z}_{ik})}$$

Let's say that we want to choose our **proposal in a parametric family** $(\Psi(\cdot|\kappa))_{\kappa\in\mathcal{K}}$ over \mathbb{R}^d (e.g. Gaussians).

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Problem: we need to choose n proposals $q_1, ..., q_n$ (and n is usually large in deep learning...).

Maximum likelihood for DLVMs: choosing proposals

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What would be the optimal, zero-variance choices for $q_1, ..., q_n$?

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What would be the optimal, zero-variance choices for $q_1, ..., q_n$?

$$q_i(\mathbf{z}) = p_{\theta}(\mathbf{z}|\mathbf{x}_i)$$

A solution: Amortised variational inference, all the q_i will be defined together via a neural net!

Rationale: q_i needs to depends on \mathbf{x}_i , so we'll define it as a conditional distribution parametrised by γ :

$$q_i(\mathbf{z}) = q_{\gamma}(\mathbf{z}|\mathbf{x}_i).$$

How to parametrise this conditional distribution? The key idea is that its parameters are the output of a neural net g_{γ} :

$$q_{\gamma}(\mathbf{z}|\mathbf{x}_i) = \Psi(\mathbf{z}|g_{\gamma}(\mathbf{x}_i)).$$

This neural net is called the **inference network** or **encoder**.

All of this leads to the following approximation of the likelihood

$$\ell(\boldsymbol{\theta}) pprox \sum_{i=1}^n \mathbb{E}_{\mathbf{z}_{i1},...,\mathbf{z}_{iK} \sim q_{\boldsymbol{\gamma}}(\mathbf{z}|\mathbf{x}_i)} \left[\log \frac{1}{K} \sum_{k=1}^K \frac{p_{\boldsymbol{\theta}}(\mathbf{x}_i|\mathbf{z}_{ik})p(\mathbf{z}_{ik})}{q_{\boldsymbol{\gamma}}(\mathbf{z}_{ik}|\mathbf{x}_i)} \right] = \mathcal{L}_K(\boldsymbol{\theta}, \boldsymbol{\gamma}).$$

Rather than maximising $\ell(\theta)$, we'll maximise $\mathcal{L}_K(\theta, \gamma)$ using SGD and the reparametrisation trick. But does it make sense to do that?

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It does make sense! For several reasons:

- $\mathcal{L}_K(\theta, \gamma)$ is a **lower bound of** $\ell(\theta)$ (exercise!)
- The bounds get tighter and tighter!

$$\mathcal{L}_1(\boldsymbol{\theta}, \boldsymbol{\gamma}) \leq \mathcal{L}_2(\boldsymbol{\theta}, \boldsymbol{\gamma}) \leq \ldots \leq \mathcal{L}_K(\boldsymbol{\theta}, \boldsymbol{\gamma}) \xrightarrow[K \to \infty]{} \ell(\boldsymbol{\theta}).$$

 $\mathcal{L}_K(\theta, \gamma)$ is called the **importance weighted autoencoder (IWAE)** bound, and was introduced by Burda et al. (2016).

The VAE bound of Kingma & Welling (2014) and Rezende et al. (2014) is actually $\mathcal{L}_1(\theta, \gamma)$, which is the loosest bound!

The VAE bound can be interestingly rewritten

$$\mathcal{L}_1(\boldsymbol{\theta}, \boldsymbol{\gamma}) = \ell(\boldsymbol{\theta}) - \mathsf{KL}\left(\prod_{i=1}^n q_{\boldsymbol{\gamma}}(\mathbf{z}_i|\mathbf{x}_i) \Big| \Big| \prod_{i=1}^n p_{\boldsymbol{\theta}}(\mathbf{z}_i|\mathbf{x}_i)\right).$$

which means that, for a given θ , the optimal $q_{\gamma}(\mathbf{z}_i|\mathbf{x}_i)$ will be as close as possible (in a KL sense) to the true posterior $p_{\theta}(\mathbf{z}_i|\mathbf{x}_i)$.

Concrete consequence: after training, we may interpret the $q_{\gamma}(\mathbf{z}_i|\mathbf{x}_i)$ as an (approachable) approximation of the (intractable) $p_{\theta}(\mathbf{z}_i|\mathbf{x}_i)$.

Is it still true when K > 1? Kind of, but it gets more complicated. Domke & Sheldon (2019) showed that, when $K \to \infty$, the the "closeness" is no longer in KL sense but in the sense of the χ divergence.

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 $\mathbb{P}(\mathfrak{F}|X)$

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mismatch between \mathbf{x}_i and its reconstruction (obtained by auto-encoding it)

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KL regulariser:

makes sure the encodings are not too far away from the prior. At the end of the day, a scatter plot of the encodings will kinda look like the prior.

This motivates the name variational auto-encoder, as we can see the loss as a KL-regularised auto-encoder loss!

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$$\log p_{\boldsymbol{\theta}}(\mathbf{x}|\mathbf{z}) = \frac{-D}{2} \log(2\pi) - D \log \sigma - \frac{1}{2\sigma^2} ||\mathbf{x} - \boldsymbol{\mu}_{\boldsymbol{\theta}}(\mathbf{z})||_2^2$$

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- The other terms do not depend on θ , so it makes sense to see $\log p_{\theta}(\mathbf{x}|\mathbf{z})$ as a reconstruction error.
- Note that the autoencoding process is stochastic, as it involves sampling $\mathbf{z} \sim q_{\gamma}(\mathbf{z}|\mathbf{x})$

• If we have a Bernoulli observation model $p_{\theta}(\mathbf{x}|\mathbf{z}) = \prod_{j=1}^{n} \mathcal{B}(x_j|\boldsymbol{\pi}_{\theta}(\mathbf{z})_j)$, we get

$$\log p_{\theta}(\mathbf{x}|\mathbf{z}) = -\sum_{j=1}^{d} XEnt(\mathbf{x}, \boldsymbol{\pi}_{\theta}(\mathbf{z}))$$

Cross-entropy loss

between \mathbf{x}_i and its reconstruction (obtained by auto-encoding it)

• The KL part of the VAE bound is equal to $\mathrm{KL}\left(\prod_{i=1}^n q_{\gamma}(\mathbf{z}_i|\mathbf{x}_i) \middle| \prod_{i=1}^n p_{\boldsymbol{\theta}}(\mathbf{z}_i)\right)$, which is the

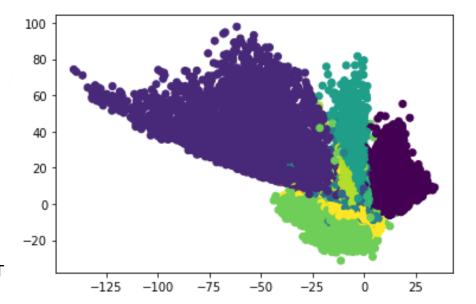
divergence between the approximate posterior (aka encoding) and the prior.

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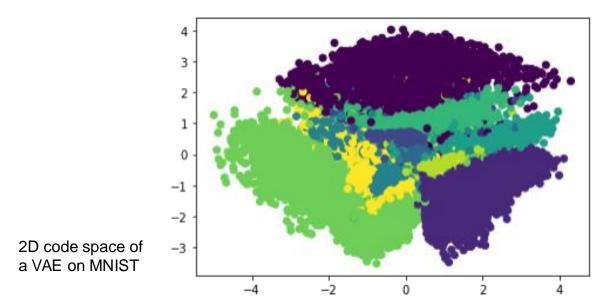
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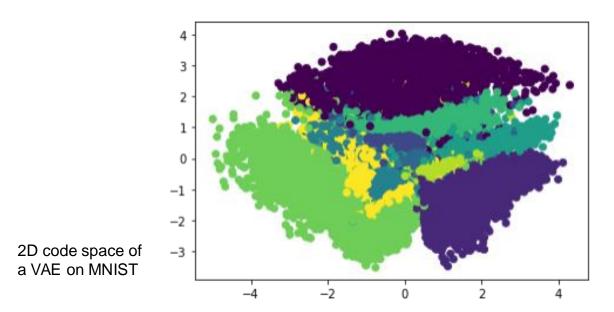
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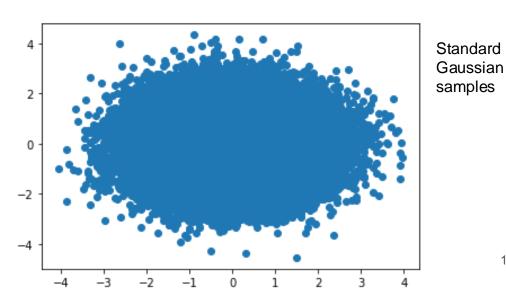


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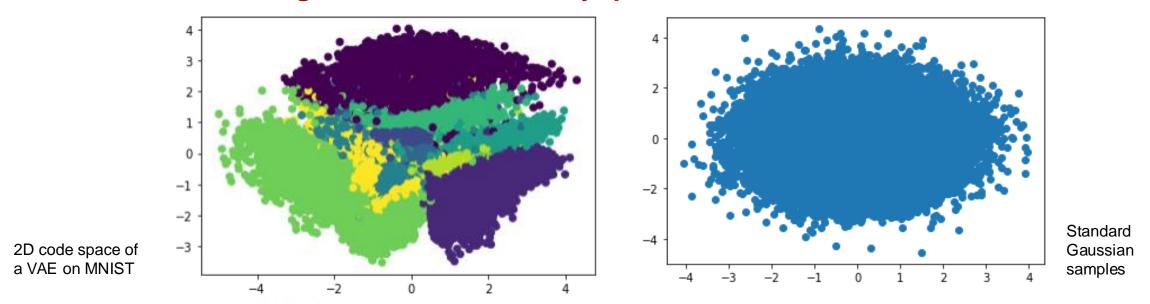
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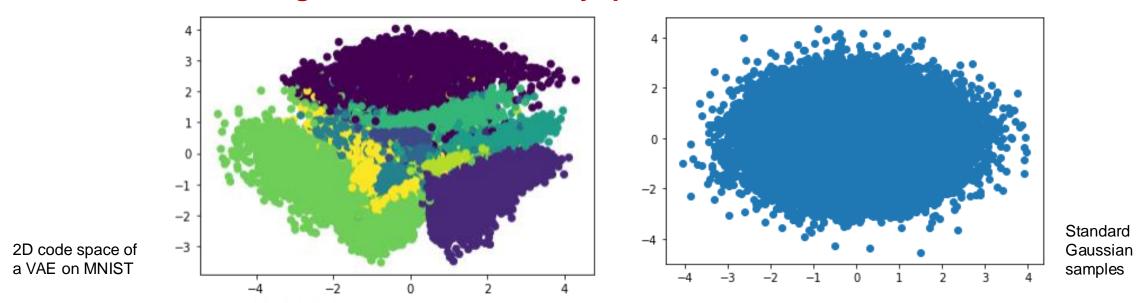
104

• So, the encodings of a VAE are actually quite Gaussian!



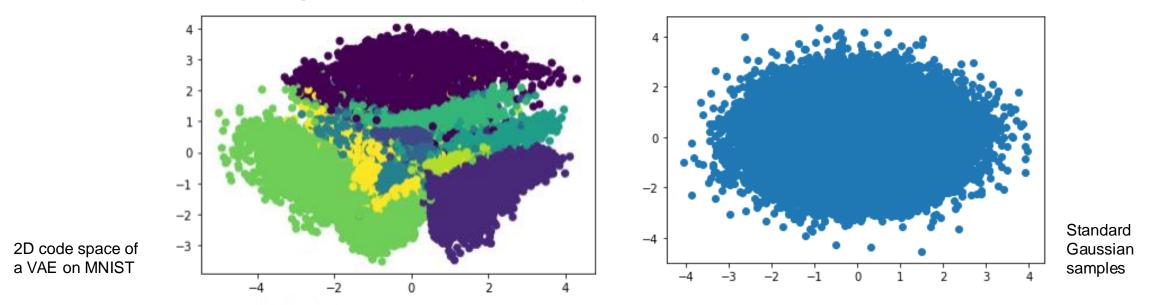
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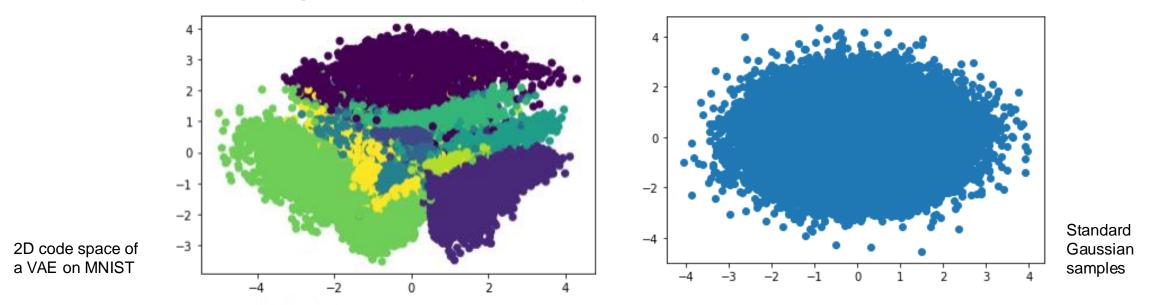
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- Is it a good or a bad thing?
 - Good: Gaussian data are well-behaved (in particular, Euclidean geometry makes sense), the latent space has no nonlinear structure
 - Bad: The latent space is not discriminative! Any idea how to simply solve this? One solution is to use a GMM prior.

More on this KL regulariser

 In the common case where both prior and approximate posterior are Gaussian, this KL regulariser is just a KL between Gaussians, which has a closed-form expression. Here is the general formula for p-variate Gaussians with full covariance

$$\mathbf{KL}(\mathcal{N}(\boldsymbol{\mu}_0, \boldsymbol{\Sigma}_0) || \mathcal{N}(\boldsymbol{\mu}_1, \boldsymbol{\Sigma}_1)) = \frac{1}{2} \left(\mathbf{tr}(\boldsymbol{\Sigma}_1^{-1} \boldsymbol{\Sigma}_0) + (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_0)^T \boldsymbol{\Sigma}_1^{-1} (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_0) + \log \frac{\det \boldsymbol{\Sigma}_1}{\det \boldsymbol{\Sigma}_0} - p \right)$$

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- It's easy to backprop through this, and this was used in the seminal VAE papers!
- Not easy to generalise beyond Gaussians...

How do we actually train VAEs?

- What have we done so far? We have created a family of lower bounds of the log-likelihood, but how do optimise them?
- Remember that the IWAE bounds are defined as

$$\mathcal{L}_{K}(\boldsymbol{\theta}, \boldsymbol{\gamma}) = \sum_{i=1}^{n} \mathbb{E}_{\mathbf{z}_{i1}, \dots, \mathbf{z}_{iK} \sim q_{\boldsymbol{\gamma}}(\mathbf{z}|\mathbf{x}_{i})} \left[\log \frac{1}{K} \sum_{k=1}^{K} \frac{p_{\boldsymbol{\theta}}(\mathbf{x}_{i}|\mathbf{z}_{ik})p(\mathbf{z}_{ik})}{q_{\boldsymbol{\gamma}}(\mathbf{z}_{ik}|\mathbf{x}_{i})} \right]$$

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- The idea is to use stochastic gradient descent (SGD, or one of its variants). What do we
 need to compute to perform SGD on an objective? Unbiased estimates of the gradients!
- As we'll see, it is doable to compute unbiased estimates of $\nabla_{\theta,\gamma} \mathcal{L}_K(\theta,\gamma)$

Unbiased IWAE gradients: looking at a more general problem

 Let's look at the problem in a more general form, we want unbiased estimates of the gradients of a function of the form

$$f(\boldsymbol{\theta}, \boldsymbol{\gamma}) = \mathbb{E}_{\mathbf{w} \sim \pi_{\boldsymbol{\gamma}}(\mathbf{w})} [g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w})]$$

In the next few slides, we will describe a few recipes to compute such estimates. This is a
task that is useful in a lot of ML contexts, e.g. reinforcement learning, explainability...
For more details, and more recipe, you may look at the following nice review

Journal of Machine Learning Research 21 (2020) 1-62

Monte Carlo Gradient Estimation in Machine Learning

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which means that we can get an unbiased estimate of $\nabla_{\theta} f(\theta, \gamma)$ by simply sampling $\mathbf{w}_1, ..., \mathbf{w}_K \sim \pi_{\gamma}(\mathbf{w})$ and then computing

$$\left(
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• So it's not that simple... Let us write the expectation:

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It makes sense to use Leibniz's rule again:

$$\nabla_{\gamma} f(\boldsymbol{\theta}, \boldsymbol{\gamma}) = \int \nabla_{\gamma} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \pi_{\boldsymbol{\gamma}}(\mathbf{w}) \right] d\mathbf{w}$$

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And now we can use the usual rule to **differentiate a product**:

$$\nabla_{\gamma} f(\boldsymbol{\theta}, \boldsymbol{\gamma}) = \int \nabla_{\gamma} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \right] \pi_{\gamma}(\mathbf{w}) d\mathbf{w} + \int g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \nabla_{\gamma} \left[\pi_{\gamma}(\mathbf{w}) \right] d\mathbf{w}$$

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 Which one of these two terms is easy to unbiasedly estimate? The first one because it is an expectation!

$$\int \nabla_{\gamma} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \right] \pi_{\boldsymbol{\gamma}}(\mathbf{w}) d\mathbf{w} \approx \frac{1}{K} \sum_{k=1}^{K} \nabla_{\gamma} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}_{k}) \right]$$

The real tricky part is what's left: the second term
$$\int g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \nabla_{\boldsymbol{\gamma}} \left[\pi_{\boldsymbol{\gamma}}(\mathbf{w}) \right] d\mathbf{w}$$

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$$\int g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \nabla_{\boldsymbol{\gamma}} \left[\pi_{\boldsymbol{\gamma}}(\mathbf{w}) \right] d\mathbf{w} = \mathbb{E}_{\mathbf{w} \sim \pi_{\boldsymbol{\gamma}}} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \nabla_{\boldsymbol{\gamma}} \log \pi_{\boldsymbol{\gamma}}(\mathbf{w}) \right]$$
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• The estimate
$$\int g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \nabla_{\boldsymbol{\gamma}} \left[\pi_{\boldsymbol{\gamma}}(\mathbf{w}) \right] d\mathbf{w} = \mathbb{E}_{\mathbf{w} \sim \pi_{\boldsymbol{\gamma}}} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \nabla_{\boldsymbol{\gamma}} \log \pi_{\boldsymbol{\gamma}}(\mathbf{w}) \right]$$
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- Can we do better? In general, not really. But in some specific cases, yes!
- In the next slide, we'll see one of such cases: the Gaussian reparametrisation trick.

- Our goal is still to estimate our tricky term $\nabla_{\gamma} \int g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \pi_{\boldsymbol{\gamma}}(\mathbf{w}) d\mathbf{w}$
- Again, the main issue is that the density depends on the parameter of interest. Can we destroy this dependence? Can we push \(\gamma\) away from the density we're integrating against?

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- In some important cases, yes! The main example is the Gaussian case $\pi_{\gamma} = \mathcal{N}(\mu_{\gamma}, \Sigma_{\gamma})$
- In this setting, it is clear that sampling $\mathbf{w} \sim \pi_{\gamma}$ can be done by computing

$$\mathbf{w} = \mu_{\gamma} + C_{\gamma} \varepsilon$$

where C_{γ} is the Cholesky decomposition of the covariance, and $\varepsilon \sim \mathcal{N}(0, \mathbf{I})$

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• But now, the only random thing is ε . This means we can rewrite our expectation as an expectation over ε

$$\nabla_{\gamma} \int g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \pi_{\gamma}(\mathbf{w}) d\mathbf{w} = \nabla_{\gamma} \int g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mu_{\gamma} + C_{\gamma} \boldsymbol{\varepsilon}) p(\boldsymbol{\varepsilon}) d\boldsymbol{\varepsilon}$$

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...and finally use Leibniz's rule

$$\nabla_{\gamma} \int g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mathbf{w}) \pi_{\gamma}(\mathbf{w}) d\mathbf{w} = \int \nabla_{\gamma} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mu_{\gamma} + C_{\gamma} \boldsymbol{\varepsilon}) \right] p(\boldsymbol{\varepsilon}) d\boldsymbol{\varepsilon}$$

$$\approx \frac{1}{K} \sum_{k=1}^{K} \nabla_{\gamma} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mu_{\gamma} + C_{\gamma} \boldsymbol{\varepsilon}_{k}) \right]$$

The estimate

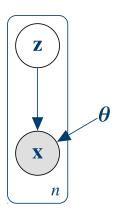
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$$\approx \frac{1}{K} \sum_{k=1}^{K} \nabla_{\gamma} \left[g(\boldsymbol{\theta}, \boldsymbol{\gamma}, \mu_{\gamma} + C_{\gamma} \boldsymbol{\varepsilon}_{k}) \right]$$

is often called the reparametrisation trick estimate. It has considerably less variance in practice than the score gradient, but can be less generally applied.

- Beyond Gaussians, this can be done for more complex distributions (Dirichlet, Student's t, GMMs), but this is not easy, in particular for discrete distributions.
- It is automatically implemented in many libraries, for instance Tensorflow Probability, or Pytorch distributions

A quick summary of VAEs/IWAEs so far

We have defined a graphical model called a deep latent variable model



 We have seen how to train this model by doing approximate maximum likelihood via amortised variational inference

 We saw that there was an important interplay between this inference technique, and various sampling techniques (importance sampling to define the bounds, various methods to estimate its gradients without bias).