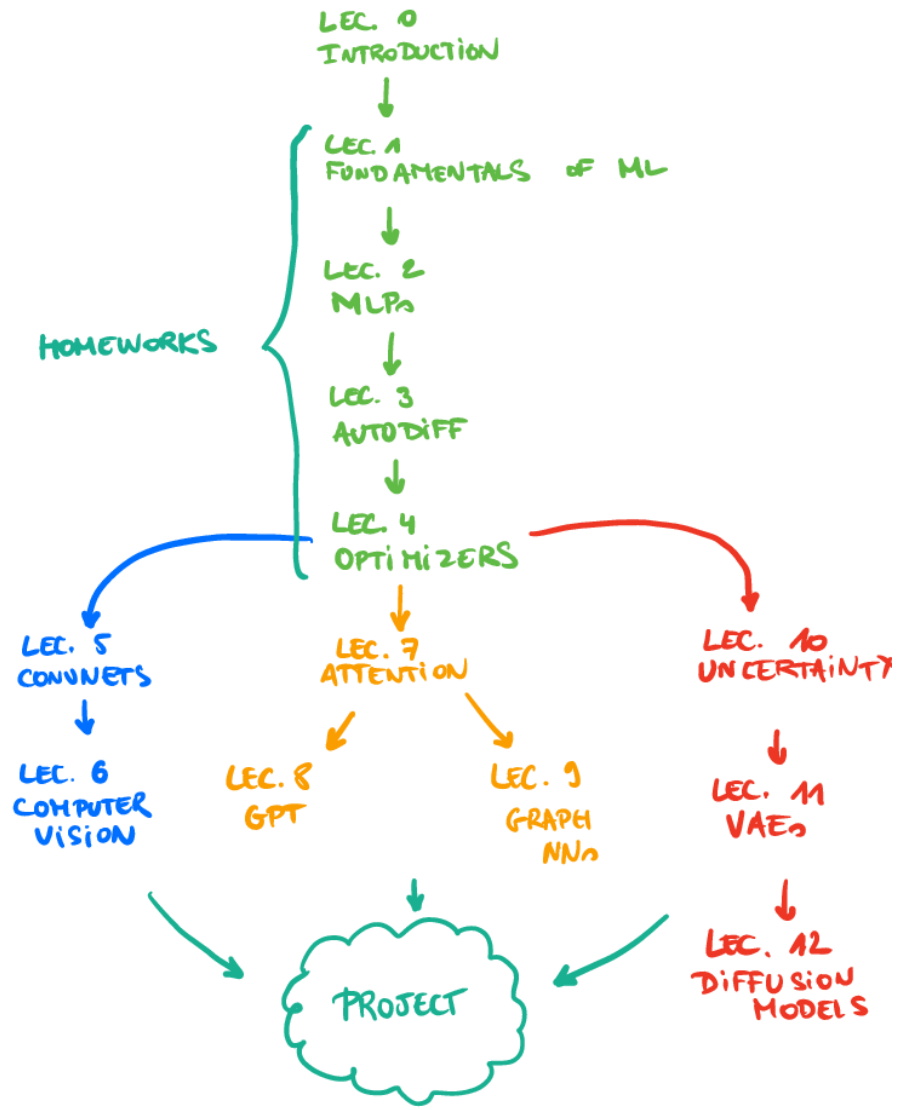


Deep Learning

Lecture 10: Uncertainty

Prof. Gilles Louppe
g.louppe@uliege.be





*"Every time a scientific paper presents a bit of data, it's accompanied by an **error bar** – a quiet but insistent reminder that no knowledge is complete or perfect. It's a **calibration of how much we trust what we think we know.**"*

Carl Sagan

Today

How to estimate uncertainty with and in neural networks?

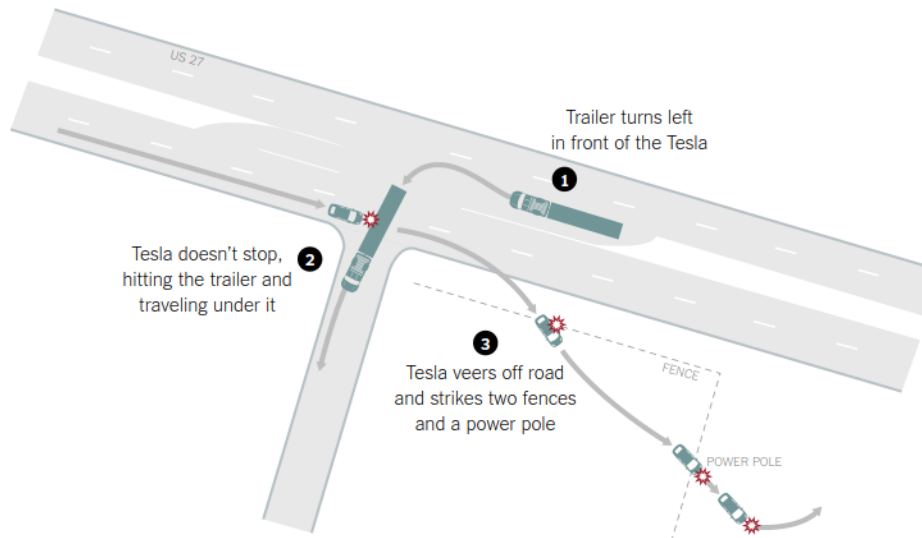
- Uncertainty
- Aleatoric uncertainty
- Epistemic uncertainty

Uncertainty

Uncertainty refers to situations where there is imperfect or unknown information. It can arise in predictions of future events, in physical measurements, or in situations where information is unknown.

Accounting for uncertainty is necessary for making optimal decisions. Not accounting for uncertainty can lead to suboptimal, wrong, or even catastrophic decisions.

Case 1. First assisted driving fatality in May 2016: Perception system mistook trailer's white side for bright sky.

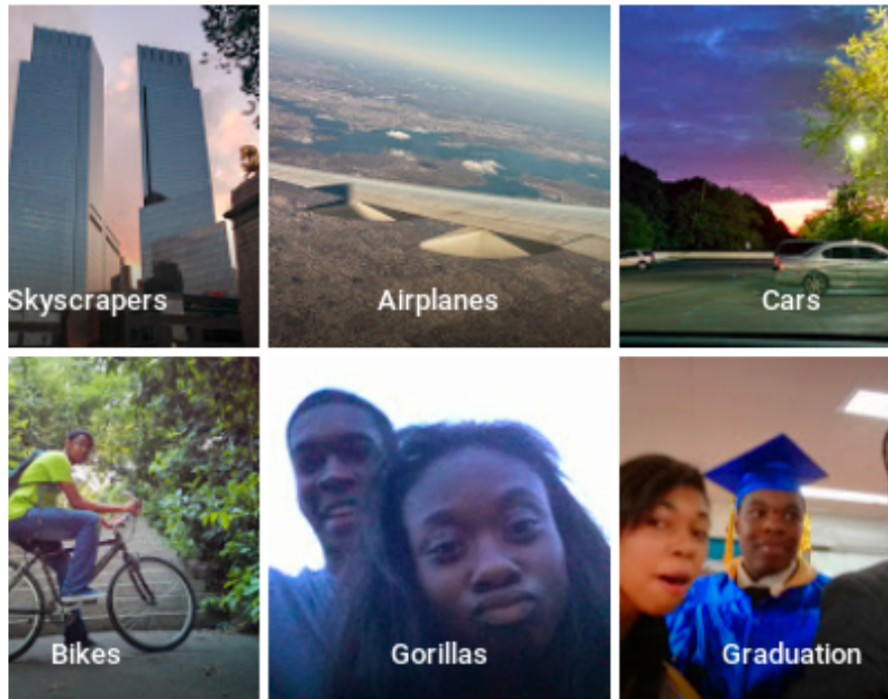




NOW
THIS

Newsflare via Reuters

**This Tesla's 'smart
summon' feature
caused it to crash into
this \$3.5M private jet**



Case 2. An image classification system erroneously identifies two African Americans as gorillas, raising concerns of racial discrimination.

The systems that made these errors were likely confident in their predictions. They did not account for uncertainty.

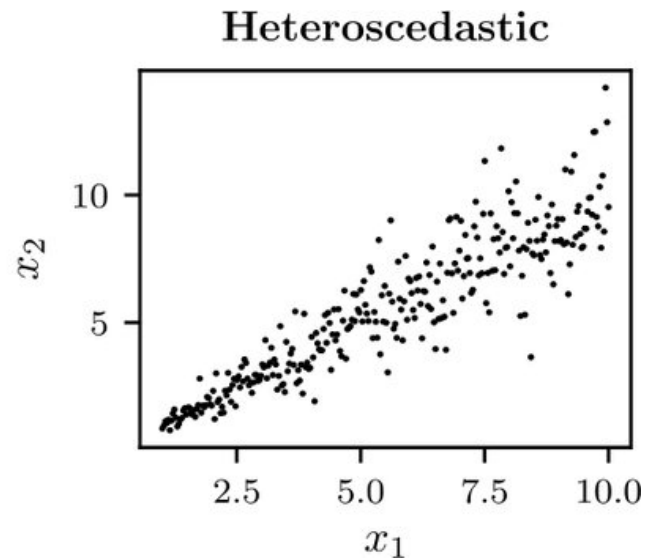
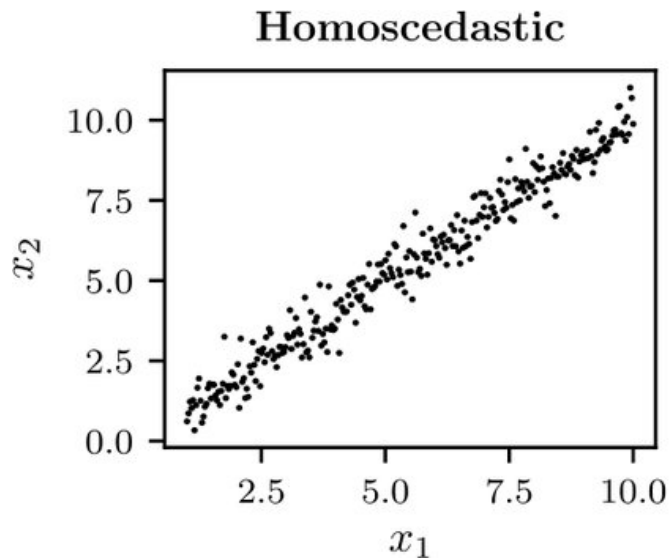
Aleatoric uncertainty

Aleatoric uncertainty refers to the uncertainty arising from the inherent stochasticity of the true data generating process. This uncertainty **cannot be reduced** with more data.

A common example is observational noise due to the limitations of the measurement devices. Collecting more data will not reduce the noise.

Assumptions about the data generating process can help in distinguishing between different types of aleatoric uncertainty:

- Homoscedastic uncertainty, which is constant across the input space.
- Heteroscedastic uncertainty, which varies across the input space.



Neural density estimation

Consider training data $(\mathbf{x}, y) \sim p(\mathbf{x}, y)$, with

- $\mathbf{x} \in \mathbb{R}^p$,
- $y \in \mathbb{R}$.

We do not wish to learn a function $\hat{y} = f(\mathbf{x})$, which would only produce point estimates.

Instead we want to learn the full conditional density

$$p(y|\mathbf{x}).$$

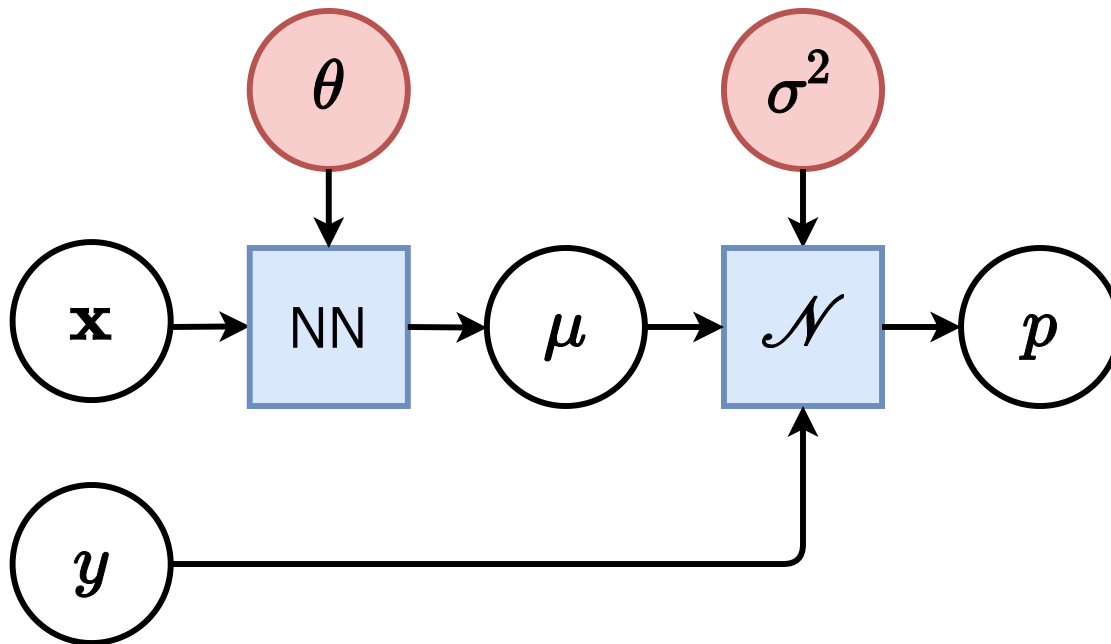
NN with Gaussian output layer

We can model aleatoric uncertainty in the output by modelling the conditional distribution as a Gaussian distribution,

$$p(y|\mathbf{x}) = \mathcal{N}(y; \mu(\mathbf{x}), \sigma^2(\mathbf{x})),$$

where $\mu(\mathbf{x})$ and $\sigma^2(\mathbf{x})$ are parametric functions to be learned, such as neural networks.

Note: The Gaussian distribution is a modelling choice. Other parametric distributions can be used.

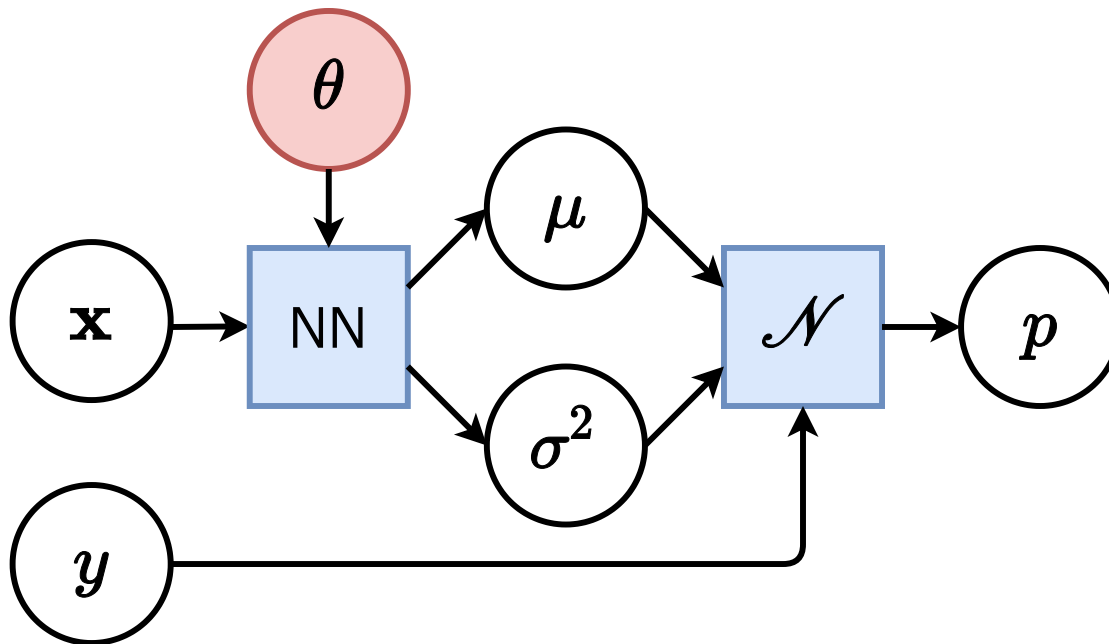


Case 1: Homoscedastic aleatoric uncertainty

We have,

$$\begin{aligned} & \arg \max_{\theta, \sigma^2} p(\mathbf{d} | \theta, \sigma^2) \\ &= \arg \max_{\theta, \sigma^2} \prod_{\mathbf{x}_i, y_i \in \mathbf{d}} p(y_i | \mathbf{x}_i, \theta, \sigma^2) \\ &= \arg \max_{\theta, \sigma^2} \prod_{\mathbf{x}_i, y_i \in \mathbf{d}} \frac{1}{\sqrt{2\pi}\sigma} \exp\left(-\frac{(y_i - \mu(\mathbf{x}_i))^2}{2\sigma^2}\right) \\ &= \arg \min_{\theta, \sigma^2} \sum_{\mathbf{x}_i, y_i \in \mathbf{d}} \frac{(y_i - \mu(\mathbf{x}_i))^2}{2\sigma^2} + \log(\sigma) + C \end{aligned}$$

What if σ^2 was fixed?



Case 2: Heteroscedastic aleatoric uncertainty

Same as for the homoscedastic case, except that that σ^2 is now a function of \mathbf{x}_i :

$$\begin{aligned} & \arg \max_{\theta} p(\mathbf{d}|\theta) \\ &= \arg \max_{\theta} \prod_{\mathbf{x}_i, y_i \in \mathbf{d}} p(y_i | \mathbf{x}_i, \theta) \\ &= \arg \max_{\theta} \prod_{\mathbf{x}_i, y_i \in \mathbf{d}} \frac{1}{\sqrt{2\pi}\sigma(\mathbf{x}_i)} \exp\left(-\frac{(y_i - \mu(\mathbf{x}_i))^2}{2\sigma^2(\mathbf{x}_i)}\right) \\ &= \arg \min_{\theta} \sum_{\mathbf{x}_i, y_i \in \mathbf{d}} \frac{(y_i - \mu(\mathbf{x}_i))^2}{2\sigma^2(\mathbf{x}_i)} + \log(\sigma(\mathbf{x}_i)) + C \end{aligned}$$

What is the purpose of $2\sigma^2(\mathbf{x}_i)$? What about $\log(\sigma(\mathbf{x}_i))$?

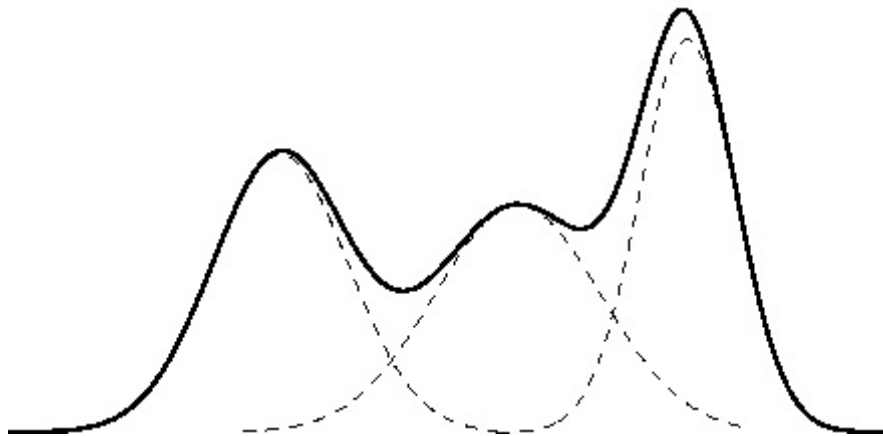
Modelling $p(y|\mathbf{x})$ as a unimodal (Gaussian) distribution can be inadequate since the conditional distribution may be **multimodal**.

Gaussian mixture model

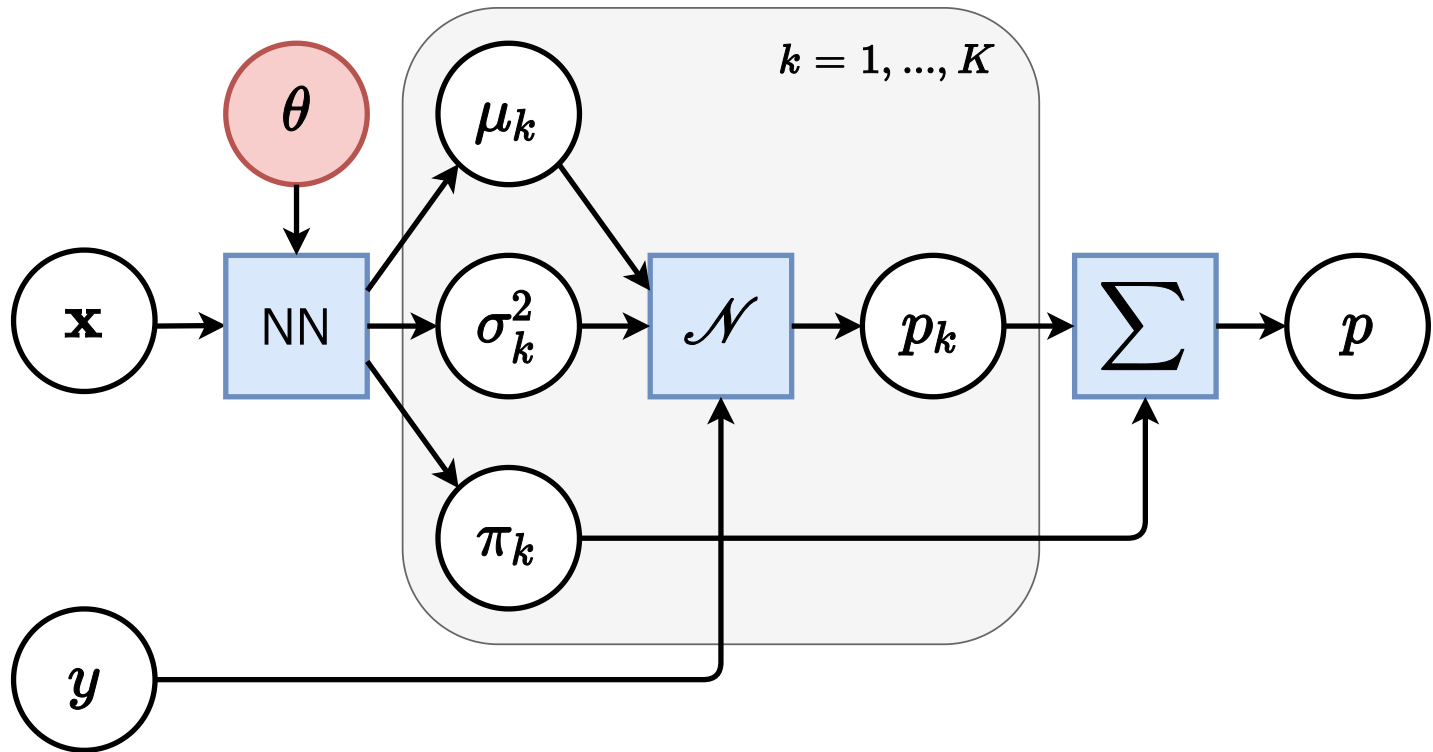
A Gaussian mixture model (GMM) defines instead $p(y|\mathbf{x})$ as a mixture of K Gaussian components,

$$p(y|\mathbf{x}) = \sum_{k=1}^K \pi_k \mathcal{N}(y; \mu_k, \sigma_k^2),$$

where $0 \leq \pi_k \leq 1$ for all k and $\sum_{k=1}^K \pi_k = 1$.



A **mixture density network** (MDN) is a neural network implementation of the Gaussian mixture model.

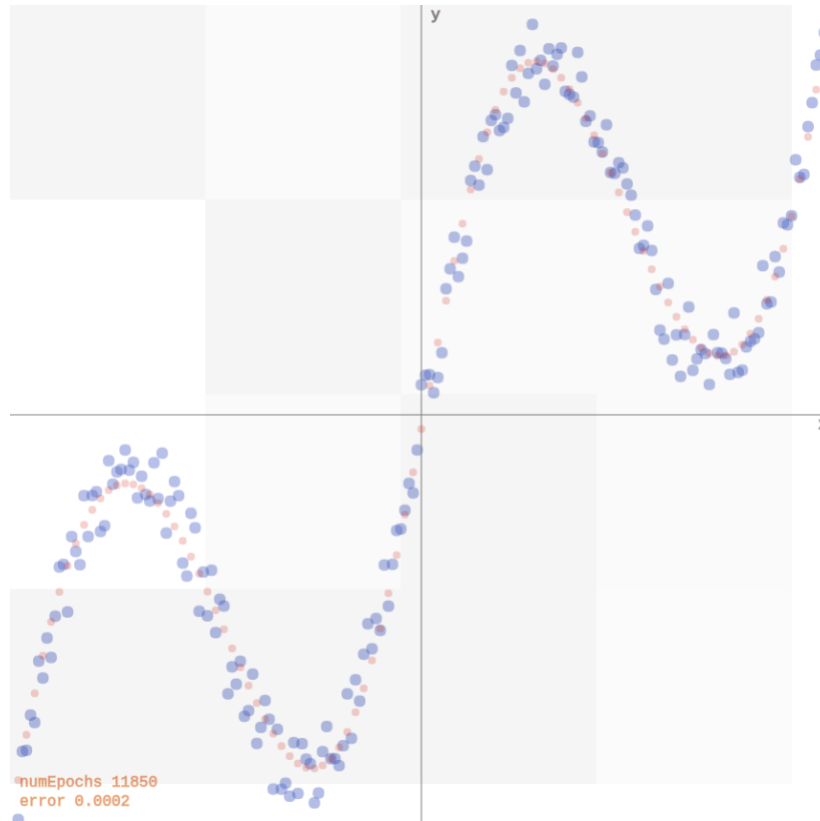


Illustration

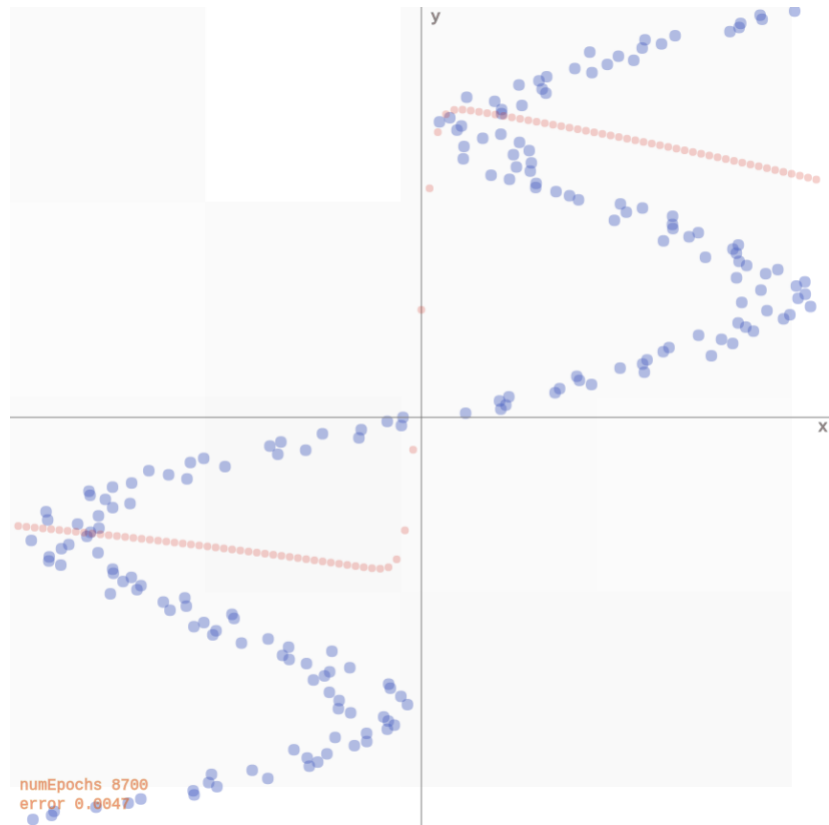
Let us consider training data generated randomly as

$$y_i = \mathbf{x}_i + 0.3 \sin(4\pi \mathbf{x}_i) + \epsilon_i$$

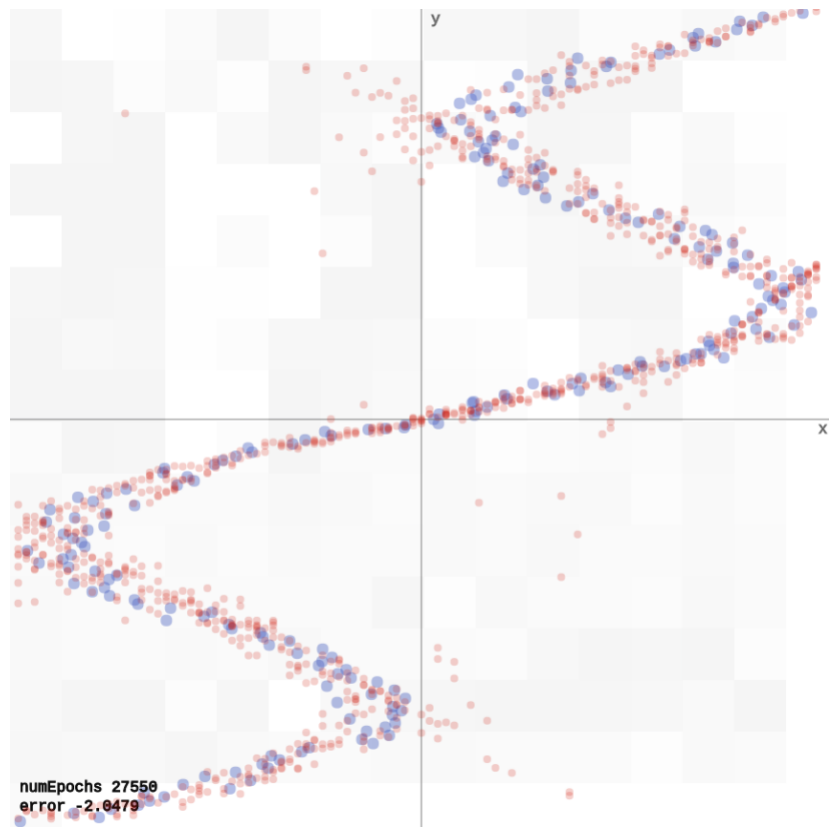
with $\epsilon_i \sim \mathcal{N}$.



The data can be fit with a 2-layer network producing point estimates for y (demo).



If we flip \mathbf{x}_i and \mathbf{y}_i , the network faces issues since for each input, there are multiple outputs that can work. It produces an average of the correct values (demo).



A mixture density network models the data correctly, as it predicts for each input a distribution for the output, rather than a point estimate ([demo](#)).

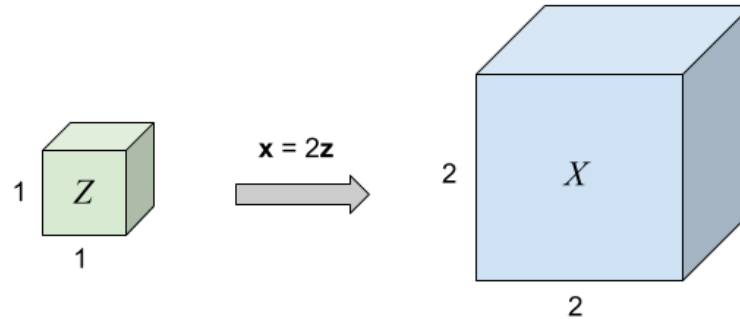
Normalizing flows

Gaussian mixture models are a flexible way to model multimodal distributions, but they are limited by the number of components K .

In practice, K must be large to model complex distributions, which makes inference difficult.

Normalizing flows are a more flexible way to model complex distributions.

Change of variables



Assume $p(\mathbf{z})$ is a uniformly distributed unit cube in \mathbb{R}^3 and $\mathbf{x} = f(\mathbf{z}) = 2\mathbf{z}$. Since the total probability mass must be conserved,

$$p(\mathbf{x}) = p(\mathbf{x} = f(\mathbf{z})) = p(\mathbf{z}) \frac{V_{\mathbf{z}}}{V_{\mathbf{x}}} = p(\mathbf{z}) \frac{1}{8},$$

where $\frac{1}{8} = \left| \det \begin{pmatrix} 2 & 0 & 0 \\ 0 & 2 & 0 \\ 0 & 0 & 2 \end{pmatrix} \right|^{-1}$ represents the inverse determinant of the linear transformation f .

What if f is non-linear?

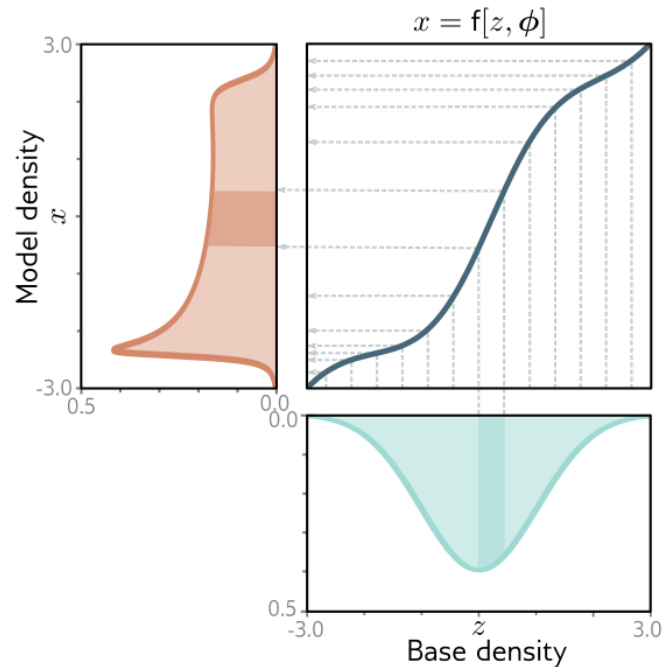


Figure 16.2 Transforming distributions. The base density (cyan, bottom) passes through a function (blue curve, top right) to create the model density (orange, left). Consider dividing the base density into equal intervals (gray vertical lines). The probability mass between adjacent lines must remain the same after transformation. The cyan-shaded region passes through a part of the function where the gradient is larger than one, so this region is stretched. Consequently, the height of the orange-shaded region must be lower so that it retains the same area as the cyan-shaded region. In other places (e.g., $z = -2$), the gradient is less than one, and the model density increases relative to the base density.

Change of variables theorem

If f is non-linear,

- the Jacobian $J_f(\mathbf{z})$ of $\mathbf{x} = f(\mathbf{z})$ represents the infinitesimal linear transformation in the neighborhood of \mathbf{z} ;
- if the function is a bijective map, then the mass must be conserved locally.

Therefore, the local change of density yields

$$p(\mathbf{x} = f(\mathbf{z})) = p(\mathbf{z}) |\det J_f(\mathbf{z})|^{-1}.$$

Similarly, for $g = f^{-1}$, we have

$$p(\mathbf{x}) = p(\mathbf{z} = g(\mathbf{x})) |\det J_g(\mathbf{x})|.$$

Example: coupling layers

Assume $\mathbf{z} = (\mathbf{z}_a, \mathbf{z}_b)$ and $\mathbf{x} = (\mathbf{x}_a, \mathbf{x}_b)$. Then,

- Forward mapping $\mathbf{x} = f(\mathbf{z})$:

$$\mathbf{x}_a = \mathbf{z}_a, \quad \mathbf{x}_b = \mathbf{z}_b \odot \exp(s(\mathbf{z}_a)) + t(\mathbf{z}_a),$$

- Inverse mapping $\mathbf{z} = g(\mathbf{x})$:

$$\mathbf{z}_a = \mathbf{x}_a, \quad \mathbf{z}_b = (\mathbf{x}_b - t(\mathbf{x}_a)) \odot \exp(-s(\mathbf{x}_a)),$$

where s and t are arbitrary neural networks.

For $\mathbf{x} = (\mathbf{x}_a, \mathbf{x}_b)$, the log-likelihood is

$$\log p(\mathbf{x}) = \log p(\mathbf{z}) |\det J_f(\mathbf{z})|^{-1}$$

where the Jacobian $J_f(\mathbf{z}) = \frac{\partial \mathbf{x}}{\partial \mathbf{z}}$ is a lower triangular matrix

$$\begin{pmatrix} \mathbf{I} & 0 \\ \frac{\partial \mathbf{x}_b}{\partial \mathbf{z}_a} & \text{diag}(\exp(s(\mathbf{z}_a))) \end{pmatrix},$$

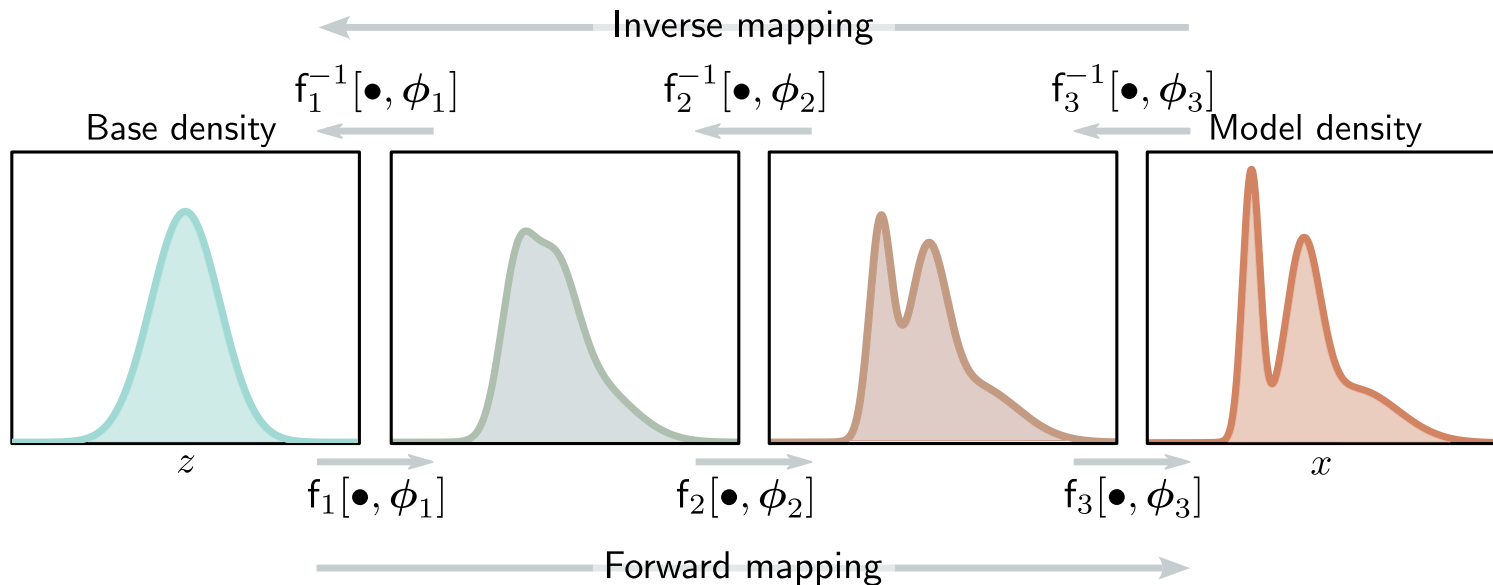
such that $|\det J_f(\mathbf{z})| = \prod_i \exp(s(\mathbf{z}_a)_i) = \exp(\sum_i s(\mathbf{z}_a)_i)$.

Therefore, the log-likelihood is

$$\log p(\mathbf{x}) = \log p(\mathbf{z}) - \sum_i s(\mathbf{z}_a)_i$$

Normalizing flows

A normalizing flow is a change of variable f that transforms a base distribution $p(\mathbf{z})$ into $p(\mathbf{x})$ through a discrete sequence of invertible transformations.



Formally,

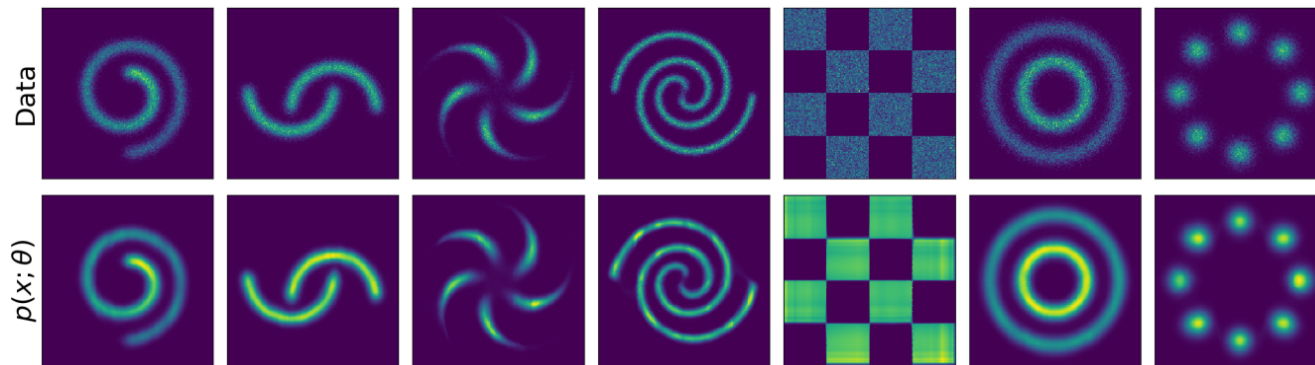
$$\mathbf{z}_0 \sim p(\mathbf{z})$$

$$\mathbf{z}_k = f_k(\mathbf{z}_{k-1}), \quad k = 1, \dots, K$$

$$\mathbf{x} = \mathbf{z}_K = f_K \circ \dots \circ f_1(\mathbf{z}_0).$$

The change of variable theorem yields

$$\log p(\mathbf{x}) = \log p(\mathbf{z}_0) - \sum_{k=1}^K \log |\det J_{f_k}(\mathbf{z}_{k-1})|.$$



Normalizing flows can fit complex multimodal discontinuous densities.

Conditional normalizing flows

Normalizing flows can also estimate densities $p(\mathbf{x}|\mathbf{c})$ conditioned on a context \mathbf{c}

- Transformations are made conditional by taking \mathbf{c} as an additional input. For example, in a coupling layer, the networks can be upgraded to $s(\mathbf{z}, \mathbf{c})$ and $t(\mathbf{z}, \mathbf{c})$.
- Optionally, the base distribution $p(\mathbf{z})$ can also be made conditional on \mathbf{c} .

(Accordingly, aleatoric uncertainty of some output \mathbf{y} conditioned on an input \mathbf{x} can be modelled by a conditional normalizing flow $p(\mathbf{y}|\mathbf{x})$ where the context \mathbf{c} is the input \mathbf{x} .)

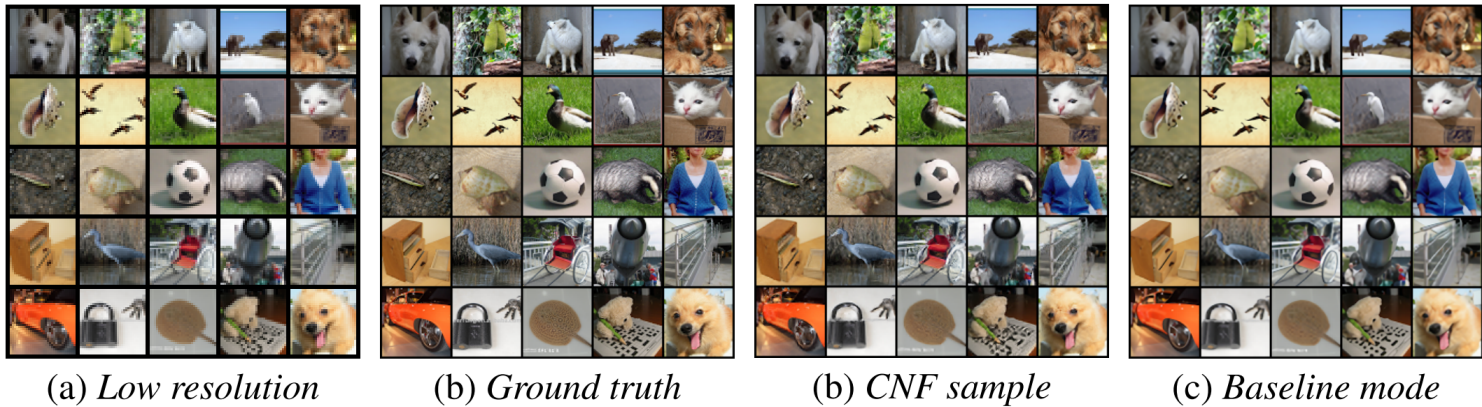


Figure 2: Super resolution results on the Imagenet64 test data. Samples are taken from the CNF $x_{hr} \sim p(x_{hr}|x_{lr})$ and the mode is visualized for the factorized baseline model. *Best viewed electronically.*

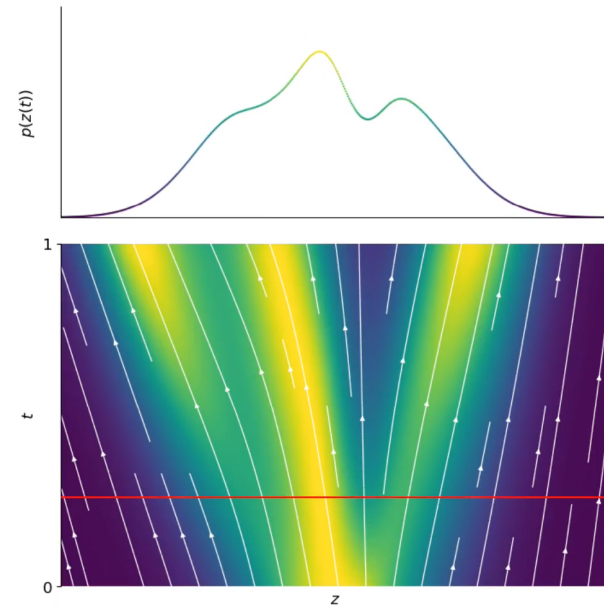
Continuous-time normalizing flows

Replace the discrete sequence of transformations with a neural ODE with reversible dynamics such that

$$\mathbf{z}_0 \sim p(\mathbf{z})$$

$$\frac{d\mathbf{z}(t)}{dt} = f(\mathbf{z}(t), t, \theta)$$

$$\mathbf{x} = \mathbf{z}(1) = \mathbf{z}_0 + \int_0^1 f(\mathbf{z}(t), t) dt.$$



The instantaneous change of variable yields

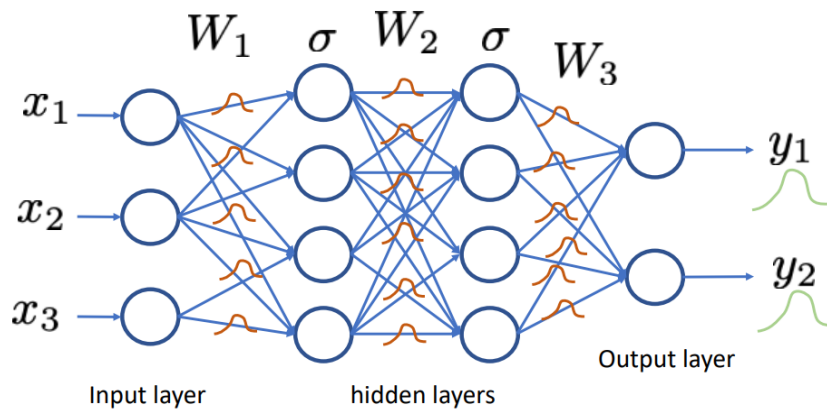
$$\log p(\mathbf{x}) = \log p(\mathbf{z}(0)) - \int_0^1 \text{Tr} \left(\frac{\partial f(\mathbf{z}(t), t, \theta)}{\partial \mathbf{z}(t)} \right) dt.$$

Epistemic uncertainty

Epistemic uncertainty accounts for uncertainty in the model or in its parameters. It captures our ignorance about which model can best explain the collected data. It **can be explained away** given enough data.

Bayesian neural networks

To capture epistemic uncertainty in a neural network, we model our ignorance with a prior distribution $p(\omega)$ over its weights and estimate the posterior distribution $p(\omega|\mathbf{d})$ given the training set \mathbf{d} .



The prior predictive distribution at \mathbf{x} is given by integrating over all possible weight configurations,

$$p(y|\mathbf{x}) = \int p(y|\mathbf{x}, \omega)p(\omega)d\omega.$$

Given training data $\mathbf{d} = \{(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)\}$ a Bayesian update results in the posterior

$$p(\omega|\mathbf{d}) = \frac{p(\mathbf{d}|\omega)p(\omega)}{p(\mathbf{d})}$$

where the likelihood $p(\mathbf{d}|\omega) = \prod_i p(y_i|\mathbf{x}_i, \omega)$.

The posterior predictive distribution is then given by

$$p(y|\mathbf{x}, \mathbf{d}) = \int p(y|\mathbf{x}, \omega)p(\omega|\mathbf{d})d\omega.$$

Bayesian neural networks are easy to formulate, but notoriously **difficult** to perform inference in.

$p(\mathbf{d})$ is intractable to evaluate, which results in the posterior $p(\omega|\mathbf{d})$ not being tractable either.

Therefore, we must rely on approximations.

Variational inference

Variational inference can be used for building an approximation $q(\omega; \nu)$ of the posterior $p(\omega|\mathbf{d})$.

We can show that minimizing

$$\text{KL}(q(\omega; \nu) || p(\omega|\mathbf{d}))$$

with respect to the variational parameters ν , is identical to maximizing the evidence lower bound objective (ELBO)

$$\text{ELBO}(\nu) = \mathbb{E}_{q(\omega; \nu)} [\log p(\mathbf{d}|\omega)] - \text{KL}(q(\omega; \nu) || p(\omega)).$$

The integral in the ELBO is not tractable for almost all q , but it can be maximized with stochastic gradient ascent:

1. Sample $\hat{\omega} \sim q(\omega; \nu)$.
2. Do one step of maximization with respect to ν on

$$\hat{L}(\nu) = \log p(\mathbf{d}|\hat{\omega}) - \log \frac{q(\hat{\omega}; \nu)}{p(\hat{\omega})}$$

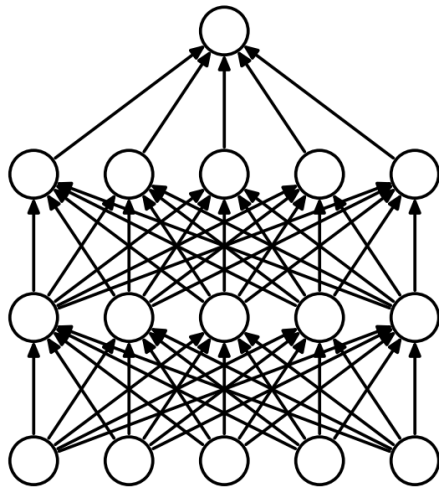
In the context of Bayesian neural networks, this procedure is also known as **Bayes by backprop** (Blundell et al, 2015).

Dropout

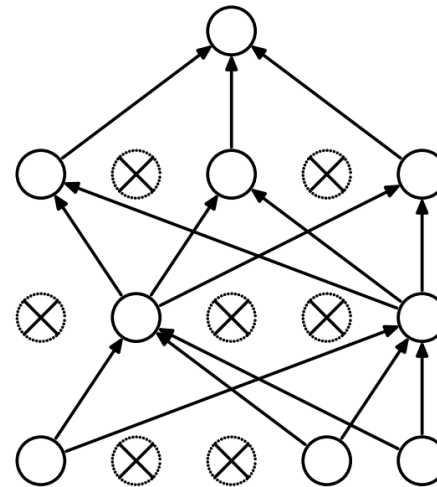
Dropout is an empirical technique that was first proposed to avoid overfitting in neural networks.

At each training step:

- Remove each node in the network with a probability p .
- Update the weights of the remaining nodes with backpropagation.



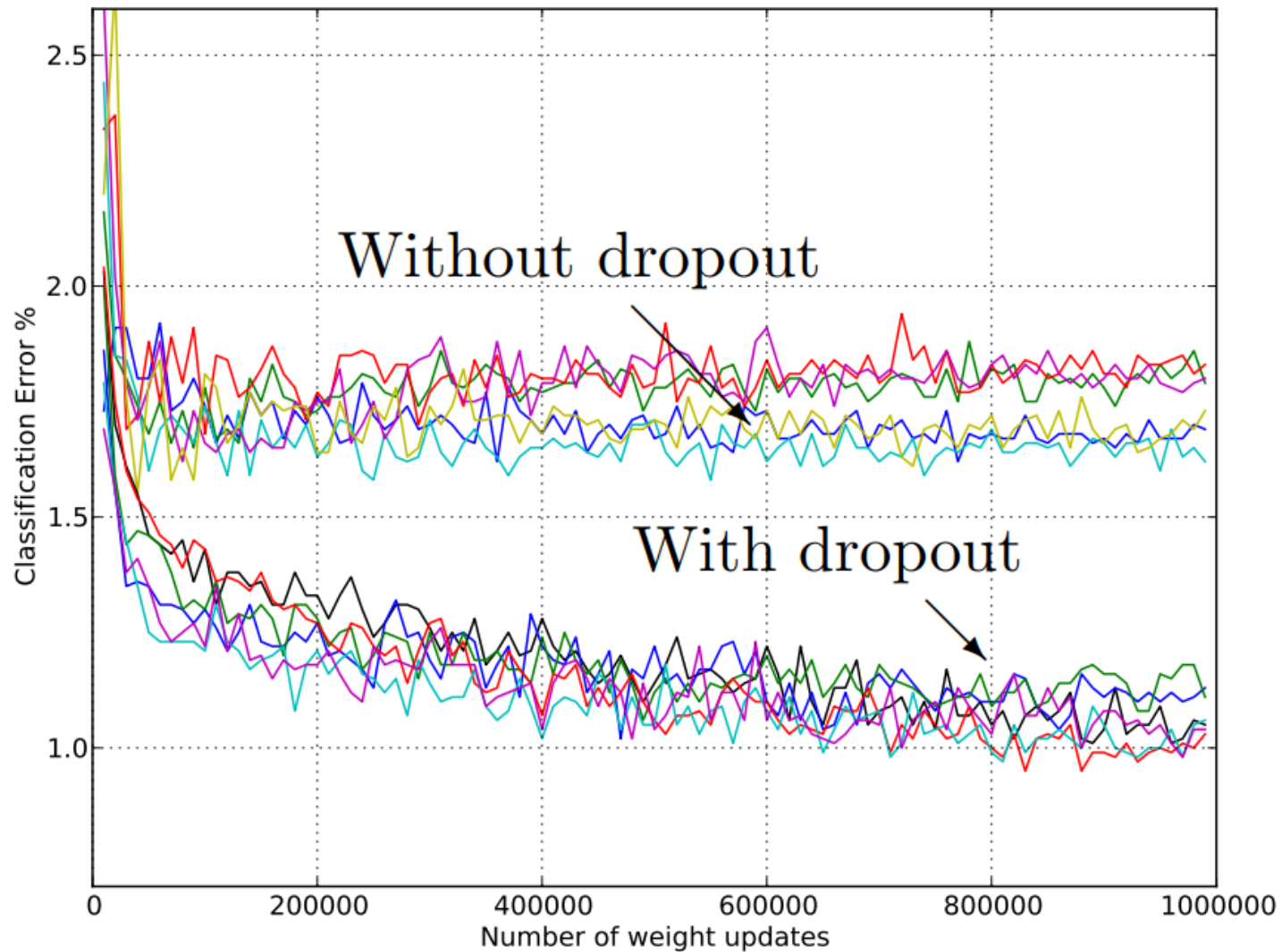
(a) Standard Neural Net



(b) After applying dropout.

At **test time**, either:

- Make predictions using the trained network **without** dropout but rescaling the weights by the dropout probability p (fast and standard).
- Sample T neural networks using dropout and average their predictions (slower but better principled).



Why does dropout work?

- It makes the learned weights of a node less sensitive to the weights of the other nodes.
- This forces the network to learn several independent representations of the patterns and thus decreases overfitting.
- It approximates **Bayesian model averaging**.

Dropout does variational inference

What variational family q would correspond to dropout?

- Let us split the weights ω per layer, $\omega = \{\mathbf{W}_1, \dots, \mathbf{W}_L\}$, where \mathbf{W}_i is further split per unit $\mathbf{W}_i = \{\mathbf{w}_{i,1}, \dots, \mathbf{w}_{i,q_i}\}$.
- Variational parameters ν are split similarly into $\nu = \{\mathbf{M}_1, \dots, \mathbf{M}_L\}$, with $\mathbf{M}_i = \{\mathbf{m}_{i,1}, \dots, \mathbf{m}_{i,q_i}\}$.
- Then, the proposed $q(\omega; \nu)$ is defined as follows:

$$q(\omega; \nu) = \prod_{i=1}^L q(\mathbf{W}_i; \mathbf{M}_i)$$
$$q(\mathbf{W}_i; \mathbf{M}_i) = \prod_{k=1}^{q_i} q(\mathbf{w}_{i,k}; \mathbf{m}_{i,k})$$
$$q(\mathbf{w}_{i,k}; \mathbf{m}_{i,k}) = p\delta_0(\mathbf{w}_{i,k}) + (1-p)\delta_{\mathbf{m}_{i,k}}(\mathbf{w}_{i,k})$$

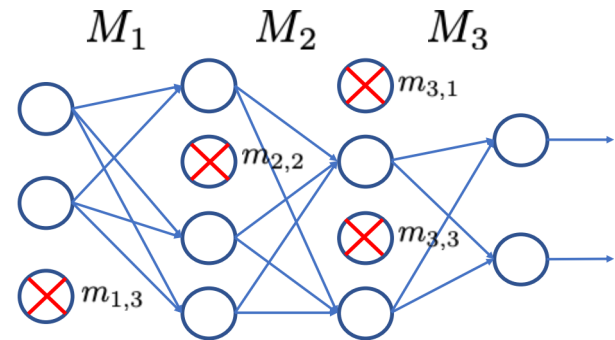
where $\delta_a(\mathbf{x})$ denotes a (multivariate) Dirac distribution centered at \mathbf{a} .

Given the previous definition for q , sampling parameters $\hat{\omega} = \{\hat{\mathbf{W}}_1, \dots, \hat{\mathbf{W}}_L\}$ is done as follows:

- Draw binary $z_{i,k} \sim \text{Bernoulli}(1 - p)$ for each layer i and unit k .
- Compute $\hat{\mathbf{W}}_i = \mathbf{M}_i \text{diag}([z_{i,k}]_{k=1}^{q_{i-1}})$, where \mathbf{M}_i denotes a matrix composed of the columns $\mathbf{m}_{i,k}$.

That is, $\hat{\mathbf{W}}_i$ are obtained by setting columns of \mathbf{M}_i to zero with probability p .

This is **strictly equivalent to dropout**, i.e. removing units from the network with probability p .



Therefore, one step of stochastic gradient descent on the ELBO becomes:

1. Sample $\hat{\omega} \sim q(\omega; \nu) \Leftrightarrow$ Randomly set units of the network to zero \Leftrightarrow Dropout.
2. Do one step of maximization with respect to $\nu = \{\mathbf{M}_i\}$ on

$$\hat{L}(\nu) = \log p(\mathbf{d}|\hat{\omega}) - \text{KL}(q(\omega; \nu)||p(\omega)).$$

Maximizing $\hat{L}(\nu)$ is equivalent to minimizing

$$-\hat{L}(\nu) = -\log p(\mathbf{d}|\hat{\omega}) + \text{KL}(q(\omega; \nu)||p(\omega))$$

This is also equivalent to one minimization step of a standard classification or regression objective:

- The first term is the typical objective (such as the cross-entropy).
- The second term forces q to remain close to the prior $p(\omega)$.
 - If $p(\omega)$ is Gaussian, minimizing the **KL** is equivalent to ℓ_2 regularization.
 - If $p(\omega)$ is Laplacian, minimizing the **KL** is equivalent to ℓ_1 regularization.

Conversely, this shows that when **training a network with dropout** with a standard classification or regression objective, one **is actually implicitly doing variational inference** to match the posterior distribution of the weights.

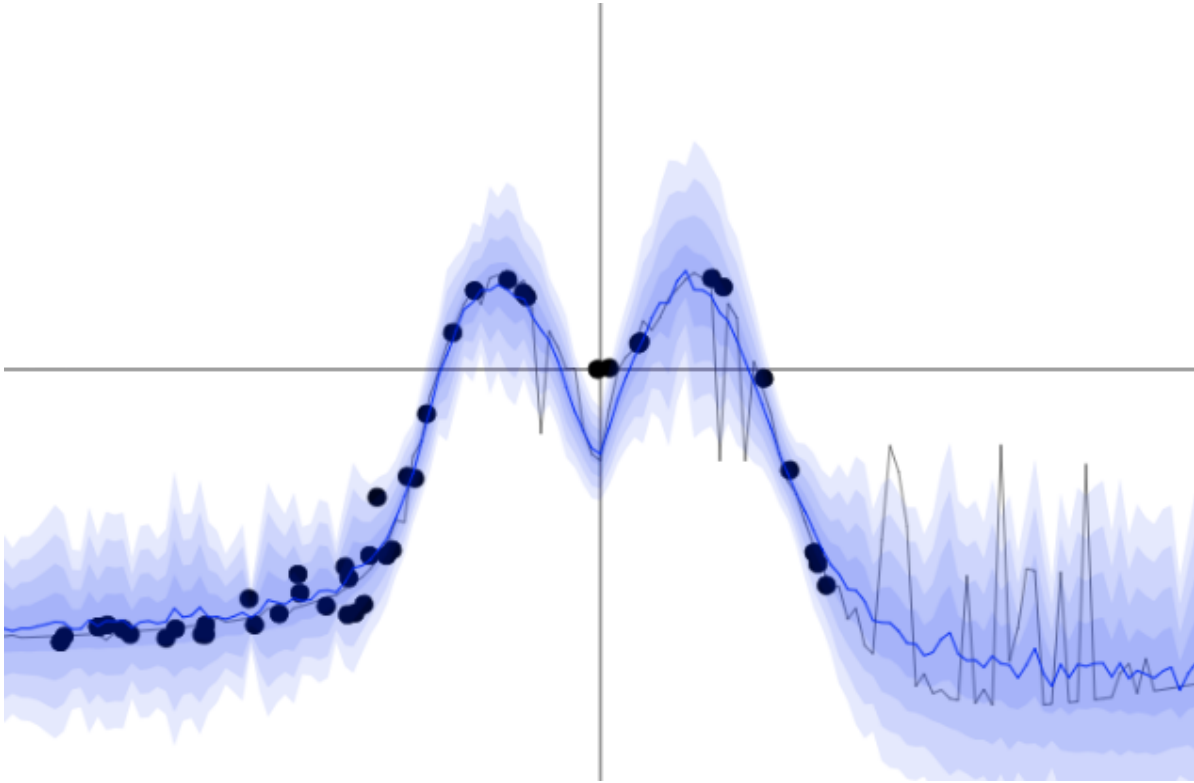
Uncertainty estimates from dropout

Proper uncertainty estimates at \mathbf{x} , accounting for both the aleatoric and epistemic uncertainties, can be obtained in a principled way using Monte-Carlo integration:

- Draw T sets of network parameters $\hat{\omega}_t$ from $q(\omega; \nu)$.
- Compute the predictions for the T networks, $\{f(\mathbf{x}; \hat{\omega}_t)\}_{t=1}^T$.
- Approximate the predictive mean and variance as

$$\mathbb{E}_{p(y|\mathbf{x},\mathbf{d})} [y] \approx \frac{1}{T} \sum_{t=1}^T f(\mathbf{x}; \hat{\omega}_t)$$
$$\mathbb{V}_{p(y|\mathbf{x},\mathbf{d})} [y] \approx \sigma^2 + \frac{1}{T} \sum_{t=1}^T f(\mathbf{x}; \hat{\omega}_t)^2 - \hat{\mathbb{E}} [y]^2 ,$$

where σ^2 is the assumed level of noise in the observational model.



(demo)

Pixel-wise depth regression

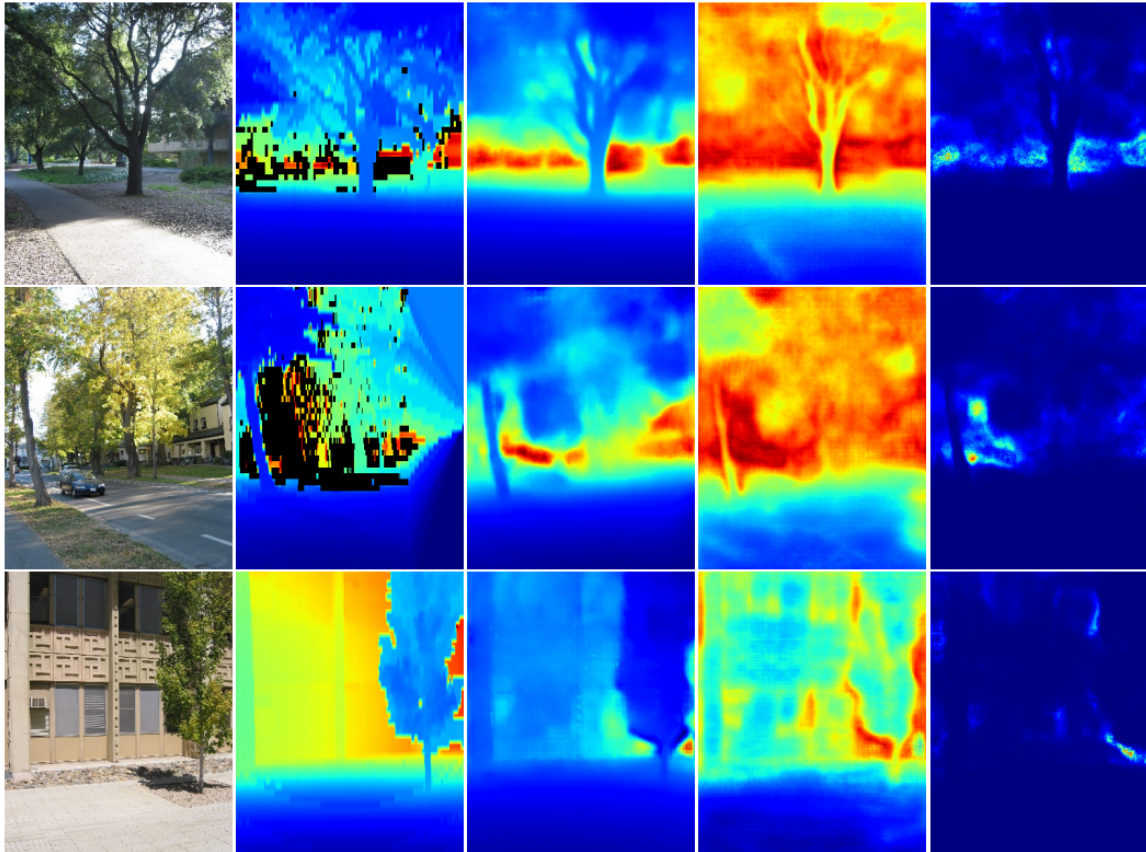


Figure 6: Qualitative results on the Make3D depth regression dataset. Left to right: input image, ground truth, depth prediction, aleatoric uncertainty, epistemic uncertainty. Make3D does not provide labels for depth greater than 70m, therefore these distances dominate the epistemic uncertainty signal. Aleatoric uncertainty is prevalent around depth edges or distant points.

