## Multivariate analysis

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- Quadratic and linear discriminants
- Support vector machines
- Kernel density estimation
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- Decision trees
(5) Summary


## Typical problems in HEP

- Classification of objects
- separate real and fake leptons/jets/etc.
- Signal enhancement relative to background
- Regression: best estimation of a parameter
- lepton energy, $\mathbb{E}_{\mathrm{T}}$ value, invariant mass, etc.


## Discrimination of signal from background in HEP

- Event level (Higgs searches, ...)
- Cone level (tau-vs-jet reconstruction, ...)
- Lifetime and flavour tagging ( $b$-tagging, )
- Track level (particle identification, ...)
- Cell level (energy deposit from hard scatter/pileup/noise, ...)


## Input information from various sources

- Kinematic variables (masses, momenta, decay angles, ... )
- Event properties (jet multiplicity, sum of charges, brightness ...)
- Event shape (sphericity, aplanarity, ...)
- Detector response (silicon hits, $d E / d x$, Cherenkov angle, shower profiles, muon hits, ...)


## Most data are (highly) multidimensional

- Use dependencies between $x=\left\{x_{1}, \cdots, x_{n}\right\}$ discriminating variables
- Approximate this $n$-dimensional space with a function $f(x)$ capturing the essential features
- $f$ is a multivariate discriminant
- For most of these lectures, use binary classification:
- an object belongs to one class (e.g. signal) if $f(x)>q$, where $q$ is some threshold,
- and to another class (e.g. background) if $f(x) \leq q$


## For simplicity: 1-dimension case

- Where to place a cut $x_{0}$ on variable $x$ ?

- Optimal choice: minimum misclassification cost at decision boundary $x=x_{0}$


## Optimal discrimination

## Cost of misclassification

$$
\begin{aligned}
C\left(x_{0}\right) & =C_{S} \int H\left(x_{0}-x\right) p(x, S) d x & & \text { signal loss } \\
& +C_{B} \int H\left(x-x_{0}\right) p(x, B) d x & & \text { background contamination }
\end{aligned}
$$

$C_{S}=$ cost of misclassifying signal as background $C_{B}=$ cost of misclassifying background as signal


- $H(x)$ : Heaviside step function
- $H(x)=1$ if $x>0$, 0 otherwise
- Optimal choice: when cost function $C$ is minimum


## Minimising the cost

- Minimise
$C\left(x_{0}\right)=C_{S} \int H\left(x_{0}-x\right) p(x, S) d x+C_{B} \int H\left(x-x_{0}\right) p(x, B) d x$ with respect to the boundary $x_{0}$ :

$$
\begin{aligned}
0 & =C_{S} \int \delta\left(x_{0}-x\right) p(x, S) d x-C_{B} \int \delta\left(x-x_{0}\right) p(x, B) d x \\
& =C_{S} p\left(x_{0}, S\right)-C_{B} p\left(x_{0}, B\right)
\end{aligned}
$$

- This gives the Bayes discriminant:

$$
B D=\frac{C_{B}}{C_{S}}=\frac{p\left(x_{0}, S\right)}{p\left(x_{0}, B\right)}=\frac{p\left(x_{0} \mid S\right) p(S)}{p\left(x_{0} \mid B\right) p(B)}
$$

## Probability relationships

- $p(A, B)=p(A \mid B) p(B)=p(B \mid A) p(A)$
- Bayes theorem: $p(A \mid B) p(B)=p(B \mid A) p(A)$
- $p(S \mid x)+p(B \mid x)=1$


## Optimal discrimination: Bayes limit

## Generalising to multidimensional problem

- The same holds when $x$ is an $n$-dimensional variable:

$$
B D=B \frac{p(S)}{p(B)} \quad \text { where } \quad B=\frac{p(x \mid S)}{p(x \mid B)}
$$

- $B$ is the Bayes factor, identical to the likelihood ratio when class densities $p(x \mid S)$ and $p(x \mid B)$ are independent of unknown parameters


## Bayes limit

- $p(S \mid x)=B D /(1+B D)$ is what should be achieved to minimise cost, achieving classification with the fewest mistakes
- Fixing relative cost of background contamination and signal loss $q=C_{B} /\left(C_{S}+C_{B}\right), q=p(S \mid x)$ defines decision boundary:
- signal-rich if $p(S \mid x) \geq q$
- background-rich if $p(S \mid x)<q$
- Any function that approximates conditional class probability $p(S \mid x)$ with negligible error reaches the Bayes limit


## How to construct $\mathrm{p}(\mathrm{S} \mid \mathrm{x})$ ?

- $k=p(S) / p(B)$ typically unknown
- Problem: $p(S \mid x)$ depends on $k$ !
- Solution: it's not a problem...
- Define a multivariate discriminant:

$$
D(x)=\frac{s(x)}{s(x)+b(x)}=\frac{p(x \mid S)}{p(x \mid S)+p(x \mid B)}
$$

- Now:

$$
p(S \mid x)=\frac{D(x)}{D(x)+(1-D(x)) / k}
$$

- Cutting on $D(x)$ is equivalent to cutting on $p(S \mid x)$, implying a corresponding (unknown) cut on $p(S \mid x)$


## Machine learning: learning from examples

## Several types of problems

- Classification/decision:
- signal or background
- type la supernova or not
- will pay his/her credit back on time or not
- Regression (mostly ignored in these lectures)
- Clustering (cluster analysis):
- in exploratory data mining, finding features


## Our goal

- Teach a machine to learn the discriminant $f(x)$ using examples from a training dataset
- Be careful to not learn too much the properties of the training sample
- no need to memorise the training sample
- instead, interested in getting the right answer for new events $\Rightarrow$ generalisation ability


## Machine learning and connected fields


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## Machine learning: (un)supervised learning

## Supervised learning

- Training events are labelled: $N$ examples $(x, y)_{1},(x, y)_{2}, \ldots,(x, y)_{N}$ of discriminating variables $x$ (also called feature variables) and class labels y
- The learner uses example classes to know how good it is doing


## Unsupervised learning

- Clustering: find similarities in training sample, without having predefined categories (how Amazon is recommending you books or DVDs...)
- Instead of categories, some sort of reward system. May not even "learn" anything from data, but remembers what triggers reward or punishment
- Not biased by pre-determined classes $\Rightarrow$ may discover unexpected features!


## A "giant" neural network

- At Google they trained a 9-layered NN with 1 billion connections
- trained on 10 million $200 \times 200$ pixel images from YouTube videos
- on 1000 machines ( 16000 cores) for 3 days, unsupervised learning
- Sounds big? The human brain has 100 billion $\left(10^{11}\right)$ neurons and 100 trillion $\left(10^{14}\right)$ connections...


## What it did

- It learned to recognise faces, one of the original goals
- ... but also cat faces (among the most popular things in YouTube videos) and body shapes



# Google's research on building high-level features 

## Features extracted from such images



- Results shown to be robust to
- colour
- translation
- scaling
- out-of-plane rotation


## Finding the multivariate discriminant $\mathrm{y}=\mathrm{f}(\mathrm{x})$

- Given our $N$ examples $(x, y)_{1}, \ldots,(x, y)_{N}$ we need
- a function class $\mathbb{F}=\{f(x, w)\}$ ( $w$ : parameters to be found)
- a constraint $Q(w)$ on $\mathbb{F}$
- a loss or error function $L(y, f)$, encoding what is lost if $f$ is poorly chosen in $\mathbb{F}$ (i.e., $f(x, w)$ far from the desired $y=f(x)$ )
- Cannot minimise $L$ directly (would depend on the dataset used), but rather its average over a training sample, the empirical risk:

$$
R(w)=\frac{1}{N} \sum_{i=1}^{N} L\left(y_{i}, f\left(x_{i}, w\right)\right)
$$

subject to constraint $Q(w)$, so we minimise the cost function:

$$
C(w)=R(w)+\lambda Q(w)
$$

- At the minimum of $C(w)$ we select $f\left(x, w_{*}\right)$, our estimate of $y=f(x)$


## Machine learning: choice of loss function

## Loss function in regression

- Goal: set $f(x, w)$ as close as possible to $y$
- Therefore, loss increases with difference between $f(x, w)$ and $y$
- Most widely used loss function is quadratic loss:

$$
L(y, f)=(f(x, w)-y)^{2}
$$

## Loss function in classification

- There is no "distance" between classes
- Goal: $f(x, w)$ predicts properly class $y$
- Usual loss function is one-loss or zero-one loss:

$$
L(y, f)=\mathbb{I}(f(x, w) \neq y)
$$

where indicator function $\mathbb{I}(X)=1$ if $X$ is true, 0 otherwise

## Choice of function class: training

Data generated from an unknown function with unknown noise


## Choice of function class: training

Constant least squares fit, RMSE $=0.915$


## Choice of function class: training



## Choice of function class: training

## Quadratic least squares fit, RMSE $=0.579$



## Choice of function class: training

## Cubic least squares fit, RMSE $=0.339$



## Choice of function class: training

Poly(6) least squares fit, RMSE $=0.278$


## Choice of function class: training

Poly(9) least squares fit, RMSE $=0$


## Quality of fit

- Increasing degree of polynomial increases flexibility of function
- Higher degree $\Rightarrow$ can match to more features
- If degree = \# points, polynomial passes through each point: perfect match!


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## Is it meaningful?

- It could be:
- if there is no noise or uncertainty in the measurement
- if the true distribution is indeed perfectly described by such a polynomial
- ... not impossible, but not very common...


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## Solution: testing sample

- Use independent sample to validate the result
- Expected: performance will also increase, go through a maximum and decrease again, while it keeps increasing on the training sample


## Choice of function class: testing

Data generated from an unknown function with unknown noise


## Choice of function class: testing

Const. least squares fit, training RMSE $=0.915$, test $\mathrm{RMSE}=1.067$

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## Choice of function class: testing

Linear least squares fit, training RMSE $=0.581$, test RMSE $=0.734$


## Choice of function class: testing

Quadr. least squares fit, training $\mathrm{RMSE}=0.579$, test $\mathrm{RMSE}=0.723$


[^0]
## Choice of function class: testing

Cubic least squares fit, training RMSE $=0.339$, test RMSE $=0.672$


## Choice of function class: testing

Poly(6) least squares fit, training RMSE $=0.278$, test RMSE $=0.72$


## Choice of function class: testing

Poly(9) least squares fit, training RMSE $=0$, test RMSE $=46.424$

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Training and test RMSE's for polinomial fits of different degrees


## Choice of function class

## Non-parametric fit

- Minimising the training cost (here, RMSE) does not work if the function class is not fixed in advance (e.g. fix the polynomial degree): complete loss of generalisation capability!
- But if you do not know the correct function class, you should not fix it! Dilemma...


## Capacity control and regularisation

- Trade-off between approximation error and estimation error
- Take into account sample size
- Measure (and penalise) complexity
- Use independent test sample
- In practice, no need to correctly guess the function class, but need enough flexibility in your model, balanced with complexity cost


## Multivariate discriminants

(4) Multivariate discriminants

- Random grid search
- Genetic algorithms
- Quadratic and linear discriminants
- Support vector machines
- Kernel density estimation
- Neural networks
- Bayesian neural networks
- Decision trees


## Reminder

- To solve binary classification problem with the fewest number of mistakes, sufficient to compute the multivariate discriminant:

$$
D(x)=\frac{s(x)}{s(x)+b(x)}
$$

where:

- $s(x)=p(x \mid S)$ signal density
- $b(x)=p(x \mid B)$ background density
- Cutting on $D(x)$ is equivalent to cutting on probability $p(S \mid x)$ that event with $x$ values is of class $S$


## Which approximation to choose?

- Best possible choice: cannot beat Bayes limit (but usually impossible to define)
- No single method can be proven to surpass all others in particular case
- Advisable to try several and use the best one


## Cut-based analysis and grid search

## Cut-based analysis

- Simple approach: cut on each discriminating variable
- Difficulty: how to optimise the cuts?


## Grid search



- Split each variable in $K$ values
- Apply cuts at each grid point: $x>x_{i}, y>y_{i}$
- Number of points scales with $K^{n}$ : curse of dimensionality

[^1]
## RGS



## Random grid search example

## Comparison to BNN




- Blue: 5-dim Bayesian neural network discriminant (see later)
- Points: each cut point from a 5-dim RGS calculation
- Conclusions:
- RGS can find very good criteria with high discrimination
- but it usually cannot compete with a full-blown multivariate discriminant
- and never outsmarts it


## Genetic algorithms

## Survival of the fittest

- Inspired by biological evolution
- Model: group (population) of abstract representations (genome/discriminating variables) of possible solutions (individuals/list of cuts)
- Typical processes at work in evolutionary processes:
- inheritance
- mutation
- sexual recombination (a.k.a. crossover)
- Fitness function: value representing the individual's goodness, or comparison of two individuals
- For cut optimisation:
- good background rejection and high signal efficiency
- compare individuals in each signal efficiency bin and keep those with higher background rejection


## Genetic algorithms

## Algorithm

- Better solutions more likely to be selected for mating and mutations, carrying their genetic code (cuts) from generation to generation
- Algorithm:
(1) Create initial random population (cut ensemble)
(2) Select fittest individuals
(3) Create offsprings through crossover (mix best cuts)
(9) Mutate randomly (change some cuts of some individuals)
(0) Repeat from 2 until convergence (or fixed number of generations)
- Good fitness at one generation $\Rightarrow$ average fitness in the next
- Algorithm focuses on region with higher potential improvement


## Gaussian problem

- Suppose densities $s(x)$ and $b(x)$ are multivariate Gaussians:

$$
\operatorname{Gaussian}(x \mid \mu, \Sigma)=\frac{1}{\sqrt{(2 \pi)^{n}|\Sigma|}} \exp \left(-\frac{1}{2}(x-\mu)^{\top} \Sigma^{-1}(x-\mu)\right)
$$

with vector of means $\mu$ and covariance matrix $\Sigma$

- Then Bayes factor $B(x)=s(x) / b(x)$ (or its logarithm) can be expressed explicitly:

$$
\ln B(x)=\lambda(x) \equiv \chi^{2}\left(\mu_{B}, \Sigma_{B}\right)-\chi^{2}\left(\mu_{S}, \Sigma_{S}\right)
$$

with $\chi^{2}(\mu, \Sigma)=(x-\mu)^{T} \Sigma^{-1}(x-\mu)$

- Fixed value of $\lambda(x)$ defines a quadratic hypersurface partitioning the $n$-dimensional space into signal-rich and background-rich regions
- Optimal separation if $s(x)$ and $b(x)$ are indeed multivariate Gaussians


## Quadratic discriminant

'Two moons' data


## Quadratic discriminant

2-D Gaussian fit for class 1

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## Quadratic discriminant

2-D Gaussian fit for class 2


## Quadratic discriminant

Discriminant function with Gaussian fits


## Linear discriminant

## Fisher's discriminant

- If in $\lambda(x)$ the same covariance matrix is used for each class (e.g. $\Sigma=\Sigma_{S}+\Sigma_{B}$ ) one gets Fisher's discriminant:

$$
\lambda(x)=w \cdot x \quad \text { with } \quad w \propto \Sigma^{-1}\left(\mu_{S}-\mu_{B}\right)
$$



- Optimal linear separation
- Works only if signal and background have different means!
- Optimal classifier (reaches the Bayes limit) for linearly correlated Gaussian-distributed variables


## Generalising Fisher discriminant

- Fisher discriminant: may fail completely for highly non-Gaussian densities
- But linearity is good feature $\Rightarrow$ try to keep it
- Idea: data non-separable in $n$-dim space $\mathbb{R}^{n}$, but better separated if mapped to higher dimension space $\mathbb{R}^{H}: h: x \in \mathbb{R}^{n} \rightarrow z \in \mathbb{R}^{H}$
- Use hyper-planes to partition higher $\operatorname{dim}$ space: $f(x)=w \cdot h(x)+b$
- Example: $h:\left(x_{1}, y_{2}\right) \rightarrow\left(z_{1}, z_{2}, z_{3}\right)=\left(x_{1}^{2}, \sqrt{2} x_{1} x_{2}, x_{2}^{2}\right)$



## Starting simple: separable data

- Consider separable data in $\mathbb{R}^{H}$, and three parallel hyper-planes:

$$
\begin{aligned}
w \cdot h(x)+b & =0(\text { separating hyper-plane between red and blue) } \\
w \cdot h\left(x_{1}\right)+b & =+1\left(\text { contains } h\left(x_{1}\right)\right) \\
w \cdot h\left(x_{2}\right)+b & =-1\left(\text { contains } h\left(x_{2}\right)\right)
\end{aligned}
$$

- Subtract blue from red:

$$
w \cdot\left(h\left(x_{1}\right)-h\left(x_{2}\right)\right)=2
$$

- With unit vector $\hat{w}=w /\|w\|$ : $\hat{w} \cdot\left(h\left(x_{1}\right)-h\left(x_{2}\right)\right)=2 /\|w\|=m$
- Margin $m$ is distance between red and blue planes
- Best separation: maximise margin
- $\Rightarrow$ empirical risk margin to minimise: $R(w) \propto\|w\|^{2}$


## Constraints

- When minimising $R(w)$, need to keep signal and background separated
- Label red dots $y=+1$ ("above" red plane) and blue dots $y=-1$ ("below" blue plane)
- Since:

$$
\begin{aligned}
& w \cdot h(x)+b>\quad 1 \text { for red dots } \\
& w \cdot h(x)+b<-1 \text { for blue dots }
\end{aligned}
$$

all correctly classified points will satisfy constraints:

$$
y_{i}\left(w \cdot h\left(x_{i}\right)+b\right) \geq 1, \forall i=1, \ldots, N
$$

- Using Lagrange multipliers $\alpha_{i}>0$, cost function can be written:

$$
C(w, b, \alpha)=\frac{1}{2}\|w\|^{2}-\sum_{i=1}^{N} \alpha_{i}\left[y_{i}\left(w \cdot h\left(x_{i}\right)+b\right)-1\right]
$$

## Minimisation

- Minimise cost function $C(w, b, \alpha)$ with respect to $w$ and $b$ :

$$
C(\alpha)=\sum_{i=1}^{N} \alpha_{i}-\frac{1}{2} \sum_{i=1}^{N} \sum_{j=1}^{N} \alpha_{i} \alpha_{j} y_{i} y_{j}\left(h\left(x_{i}\right) \cdot h\left(x_{j}\right)\right)
$$

- At minimum of $C(\alpha)$, only non-zero $\alpha_{i}$ correspond to points on red and blue planes: support vectors


## Kernel functions

- Issues:
- need to find $h$ mappings (potentially of infinite dimension)
- need to compute scalar products $h\left(x_{i}\right) \cdot h\left(x_{j}\right)$
- Fortunately $h\left(x_{i}\right) \cdot h\left(x_{j}\right)$ are equivalent to some kernel function $K\left(x_{i}, x_{j}\right)$ that does the mapping and the scalar product:

$$
C(\alpha)=\sum_{i=1}^{N} \alpha_{i}-\frac{1}{2} \sum_{i=1}^{N} \sum_{j=1}^{N} \alpha_{i} \alpha_{j} y_{i} y_{j} K\left(x_{i}, x_{j}\right)
$$

## Example

- $h:\left(x_{1}, y_{2}\right) \rightarrow\left(z_{1}, z_{2}, z_{3}\right)=\left(x_{1}^{2}, \sqrt{2} x_{1} x_{2}, x_{2}^{2}\right)$

$$
h(x) \cdot h(y)=\left(x_{1}^{2}, \sqrt{2} x_{1} x_{2}, x_{2}^{2}\right) \cdot\left(y_{1}^{2}, \sqrt{2} y_{1} y_{2}, y_{2}^{2}\right)
$$

$$
=(x \cdot y)^{2}
$$

$$
=K(x, y)
$$



- In reality: do not know a priori the right kernel
- $\Rightarrow$ have to test different standard kernels and use the best one


## Real life: non-separable data

- Even in infinite dimension space, data are often non-separable
- Need to relax constraints:

$$
y_{i}\left(w \cdot h\left(x_{i}\right)+b\right) \geq 1-\xi_{i}
$$

 with slack variables $\xi_{i}>0$

- $C(w, b, \alpha, \xi)$ depends on $\xi$, modified $C(\alpha, \xi)$ as well
- Values determined during minimisation


## Basic principle

- Introduced by E. Parzen in the 1960s
- Place a kernel $K(x, \mu)$ at each training point $\mu$
- Density $p(x)$ at point $x$ approximated by:

$$
p(x) \approx \hat{p}(x)=\frac{1}{N} \sum_{j=1}^{N} K\left(x, \mu_{j}\right)
$$




## Kernel density estimation (KDE)

## Choice of kernel

- Any kernel can be used
- In practice, often product of Gaussians:

$$
K(x, \mu)=\prod_{i}^{n} \operatorname{Gaussian}\left(x_{i} \mid \mu, h_{i}\right)
$$

each with bandwidth (width) $h_{i}$

## Optimal bandwidth

- Too narrow: noisy approximation
- Too wide: loose fine structure
- In principle found by minimising risk function

$$
R(\hat{p}, p)=\int(\hat{p}(x)-p(x))^{2} d x
$$

- For Gaussian densities:

$$
h=\sigma\left(\frac{4}{(n+2) N}\right)^{1 /(n+4)}
$$

- Far from optimal for non-Gaussian densities


## Kernel density estimation (KDE)

## Example

with Gaussian optimal bandwidth

with optimised bandwidth


## Kernel density estimation (KDE)

## Why does it work?

- When $N \rightarrow \infty$ :

$$
\hat{p}(x)=\int K(x, \mu) p(\mu) d \mu
$$

- $p(\mu)$ : true density of $x$
- Kernel bandwidth getting smaller with $N$, so when $N \rightarrow \infty$, $K(x, \mu) \rightarrow \delta^{n}(x-\mu)$ and $\hat{p}(x)=p(x)$
- KDE gives consistent estimate of probability density $p(x)$


## Limitations

- Choice of bandwidth non-trivial
- Difficult to model sharp structures (e.g. boundaries)
- Kernels too far apart in regions of low point density
- (both can be mitigated with adaptive bandwidth choice)
- Requires evaluation of $N$ n-dimensional kernels


## Kernel density estimation (KDE)

'Two moons' data


## Kernel density estimation (KDE)

2-D Parzen fit for class $1, h=2$.


## Kernel density estimation (KDE)

2-D Parzen fit for class $-1, h=2$.


## Kernel density estimation (KDE)

Discriminant function with Parzen fits, $h=2$.


## Kernel density estimation (KDE)



## Kernel density estimation (KDE)



## Kernel density estimation (KDE)

Discriminant function with Parzen fits, $h=0.01$


## Kernel density estimation (KDE)



## Kernel density estimation (KDE)



## Kernel density estimation (KDE)

Discriminant function with Parzen fits, $h=0.25$


## KDE: choice of bandwidth

Training and test error rates

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## Neural networks



## Brief history of artificial neural networks

- 1943: W. McCulloch and W. Pitts explore capabilities of networks of simple neurons
- 1958: F. Rosenblatt introduces perceptron (single neuron wih adjustable weights and threshold activation function)
- 1969: M. Minsky and S. Papert prove limitations of perceptron (linear separation only) and (wrongly) conjecture that multi-layered perceptrons have same limitations
$\Rightarrow$ ANN research almost abandoned in 1970s!!!
- 1986: Rumelhart, Hinton and Williams introduce "backward propagation of errors": solves multi-layered learning
- Today: will only talk about multilayer perceptron (MLP), but there are many recent advances in ANN


## Single neuron

- Remember linear separation:

$$
\lambda(x)=w \cdot x=\sum_{i=1}^{n} w_{i} x_{i}+w_{0}
$$

- Boundary at $\lambda(x)=0$
- Replace threshold boundary by sigmoid:


- $\sigma(\lambda)$ is neuron activity, $\lambda$ is activation
- Neuron behaviour completely controlled by weights $w=\left\{w_{0}, \ldots, w_{n}\right\}$
- Training: minimisation of error/loss function (quadratic deviations, entropy [maximum likelihood]), via gradient descent or stochastic approximation


## Training

- Minimise error function $E(w)$
- Gradient descent: $w^{(k+1)}=w^{(k)}-\eta \frac{d E^{(k)}}{d w}$
- $\frac{\partial E}{\partial w_{j}}=\sum_{n=1}^{N}-\left(t^{(n)}-y^{(n)}\right) x_{j}^{(n)}$ with target $t^{(n)}(0$ or 1$)$, so $t^{(n)}-y^{(n)}$ is the error on event $n$
- All events at once (batch learning):
- weights updated all at once after processing the entire training sample
- finds the actual steepest decent
- takes more time
- or one-by-one (online learning):
- speeds up learning
- useful in HEP because of redundant datasets (large Monte Carlo samples with many similar events)
- may avoid local minima with stochastic component in minimisation
- depends on the order of training events
- One epoch: going through the training data once


## Overtraining



- Diverging weights can cause overfitting
- Mitigate by:
- early stopping (after a fixed number of epochs)
- monitoring error on test sample
- regularisation, introducing a "weight decay" term:

$$
\tilde{E}(w)=E(w)+\frac{\alpha}{2} \sum_{i} w_{i}^{2}
$$

## Theorem

Let $\sigma($.$) be a non-constant, bounded, and monotone-increasing continuous$ function. Let $\mathcal{C}\left(I_{n}\right)$ denote the space of continuous functions on the $n$-dimensional hypercube. Then, for any given function $f \in \mathcal{C}\left(I_{n}\right)$ and $\varepsilon>0$ there exists an integer $M$ and sets of real constants $w_{j}, w_{i j}$ wherei $=1, \ldots, n$ and $j=1, \ldots, M$ such that

$$
y(x, w)=\sum_{j=1}^{M} w_{j} \sigma\left(\sum_{i=1}^{n} w_{i j} x_{i}+w_{0 j}\right)
$$

is an approximation of $f($.$) , that is |y(x)-f(x)|<\varepsilon$

## Interpretation

- You can approximate any continuous function to arbitrary precision with a linear combination of sigmoids
- Corollary 1: can approximate any continuous function with neurons!
- Corollary 2: a single hidden layer is enough
- Corollary 3: a linear output neuron is enough


## Multilayer perceptron: feedforward network

- Neurons organised in layers
- Output of one layer becomes input to next layer

$$
y_{k}(x, w)=\sum_{j=0}^{M} w_{k j}^{(2)} \underbrace{\sigma\left(\sum_{i=0}^{n} w_{j i}^{(1)} x_{i}\right)}_{z_{j}}
$$



## Can fit any function: examples

- 1 input (training data), 1 output
- 3 hidden neurons on one hidden layer
(C)Jan Therhaag
$\begin{array}{lll}-------z_{1} & \quad \text { output } \\ -------z_{2} & -. . . . . . . . . . . . . . . . . . . ~ t r a i n i n g ~ d a t a ~\end{array}$



## Backpropagation

- Training means minimising error function $E(w)$
- For single neuron: $\frac{d E}{d w_{k}}=(y-t) x_{k}$
- One can show that for a network:

$$
\frac{d E}{d w_{j i}}=\delta_{j} z_{i}, \text { where }
$$



- As before, weights can be regularised:
$\delta_{k}=\left(y_{k}-t_{k}\right)$ for output neurons
$\delta_{j} \propto \sum_{k} w_{k j} \delta_{k}$ otherwise
- Hence errors are propagated backwards


## Regularisation

## 10 hidden nodes



## 10 hidden nodes and $\alpha=0.04$



- Much less overfitting, better generalisation properties


## Getting confused: testing better than training?

- Train on noisy data centred on true value
- Test on no-noise data
- Testing error becomes better: during training, the NN learned the true distribution (average
 of noisy inputs)
- $\Rightarrow$ testing converges
- Example: $\sin (x) / x+\operatorname{rand}(-0.05,+0.05)$
- Of course doesn't work as well if noise is not symmetric


## Tricks of the trade

- Preprocess data:
- if relevant, provide e.g. $x / y$ instead of $x$ and $y$
- subtract the mean because the sigmoid derivative becomes negligible very fast
- normalise variances (close to 0 )
- shuffle training sample (order matters in online training)
- Initial random weights should be small to avoid saturation
- Batch/online training: depends on the problem
- Regularise weights to minimise overtraining. May also help select good variables via Automatic Relevance Determination (ARD)
- Make sure the training sample covers the full parameter space
- No rule (not even valid guestimates) about the number of hidden nodes
- A single hidden layer is enough for all purposes, but multiple hidden layers may allow for a solution with fewer parameters


## Adding a hidden layer



## Introduction

- As name says: Bayesian approach, try to infer functions $f(x)$
- Training sample $T$ of $N$ examples $(x, y)_{1},(x, y)_{2}, \ldots,(x, y)_{N}$ of discriminating variables $x$ and class labels $y$
- Each point $w$ corresponds to a function $f(x, w)$
- Assign probability density $p(w \mid T)$ to it
- If $p\left(w_{1} \mid T\right)>p\left(w_{2} \mid T\right)$, then associated function $f\left(x, w_{1}\right)$ more compatible with training data $T$ than function $f\left(x, w_{2}\right)$
- Posterior density $p(w \mid T)$ is final result of Bayesian inference
- BNN is the predictive distribution

$$
p(y \mid x, T)=\int p(y \mid x, w) p(w \mid T) d w
$$

where the function class is class of feedforward neural networks with a fixed structure (inputs, layers, hidden nodes, outputs)

## In practice

- Take the mean of the predictive distribution:

$$
\begin{aligned}
y(x) & =\int z p(z \mid x, T) d z \\
& =\int f(x, w) p(w \mid T) d w
\end{aligned}
$$

- Why? For classification $p(y \mid x, w)=f(x, w)^{y}(1-f(x, w))^{1-y}$
- for $y=1: p(y \mid x, w)=f(x, w)$
- for $y=0: p(y \mid x, w)=1-f(x, w)$
- so only $f(x, w)$ contributes to the mean
- Example usage:

$$
\begin{aligned}
f(x, w) & =\frac{1}{1+e^{-g(x, w)}} \\
g(x, w) & =b+\sum_{j=1}^{H} v_{j} \tanh \left(a_{j}+\sum_{i=1}^{n} u_{i j} x_{i}\right)
\end{aligned}
$$

## Implementation

- Scanning NN parameter space can be daunting
- Can approximate integral in $y(x)$ using Markov chain Monte Carlo method (MCMC)
- Will generate $M$ sample weights $w_{1}, \ldots, w_{M}$ from posterior density $p(w \mid T)$
- $y(x) \approx \frac{1}{M} \sum_{m=1}^{M} f\left(x, w_{m}\right)$
- Use spare subset of MCMC points to avoid correlations
- Start with "reasonable" guesses for parameters (e.g. zero-centred Gaussians)


## Example




- points: bin by bin histogram ratio
- thin curves: each $f\left(x, w_{k}\right)$
- thick curve: average, which approximates $D(x)$


## Details tomorrow

## Summary of MVA techniques

## Classifiers

| Criteria |  | Classifiers |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Cuts | Likeli- | PDERS | H-Matri | Fisher | MLP | BDT | RuleFit | SVM |
| Performance | no / linear correlations | () | () | () | - | () | () | () | () | () |
|  | nonlinear correlations | - | (2) | () | (\%) | (2) | (-) | () | (-) | (-) |
| Speed | Training | - | () | () | () | () | - | (2) | - | (2) |
|  | Response | () | () | ( $/$ / ${ }^{\text {c }}$ | (-) | () | () | () | - | $\bigcirc$ |
| Robust-ness | Overtraining | () | - | - | () | () | (2) | (2) | - | - |
|  | Weak input variables | () | () | (2) | () | () | (-) | (-) | - | (-) |
| Curse of dimensionality |  |  |  |  |  |  |  |  |  |  |
| Transparency |  | () | () | - | () | () | (2) | (2) | (2) | (2) |

## (according to TMVA authors)

- When trying to achieve optimal discrimination one can try to approximate

$$
D(x)=\frac{s(x)}{s(x)+b(x)}
$$

- Many techniques and tools exist to achieve this
- (Un)fortunately, no one method can be shown to outperform the others in all cases.
- One should try several and pick the best one for any given problem
- Multivariate techniques are at work in your everyday life without your knowning and can easily outsmart you for many tasks
- Try this to convince yourself http//mww.phit de/mousegame/index eng hitil


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[^0]:    (C) Balàzs Kégl

[^1]:    (c) Harrison Prosper

