Multivariate analysis

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Outline



Introduction

Optimal discrimination

- Bayes limit
- Multivariate discriminant

3 Machine learning

- Supervised and unsupervised learning
- Example

Multivariate discriminants

- Random grid search
- Genetic algorithms
- Quadratic and linear discriminants
- Support vector machines
- Kernel density estimation
- Neural networks
- Bayesian neural networks
- 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, ∉_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, ...)
- \bullet Cell level (energy deposit from hard scatter/pileup/noise, \ldots)



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, dE/dx, Cherenkov angle, shower profiles, muon hits, ...)

Most data are (highly) multidimensional

- Use dependencies between $x = \{x_1, \cdots, x_n\}$ 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 x₀ on variable x?





Cost of misclassification



• Optimal choice: when cost function C is minimum

Optimal discrimination: Bayes discriminant

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Minimising the cost

Minimise

 $C(x_0) = C_S \int H(x_0 - x)p(x, S)dx + C_B \int H(x - x_0)p(x, B)dx$ with respect to the boundary x_0 :

$$0 = C_{S} \int \delta(x_{0} - x) p(x, S) dx - C_{B} \int \delta(x - x_{0}) p(x, B) dx$$

= $C_{S} p(x_{0}, S) - C_{B} p(x_{0}, B)$

• This gives the Bayes discriminant:

$$BD = \frac{C_B}{C_S} = \frac{p(x_0, S)}{p(x_0, B)} = \frac{p(x_0|S)p(S)}{p(x_0|B)p(B)}$$

Probability relationships

•
$$p(A,B) = p(A|B)p(B) = p(B|A)p(A)$$

• Bayes theorem: p(A|B)p(B) = p(B|A)p(A)

•
$$p(S|x) + p(B|x) = 1$$

Optimal discrimination: Bayes limit



Generalising to multidimensional problem

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

$$BD = Brac{p(S)}{p(B)}$$
 where $B = rac{p(x|S)}{p(x|B)}$

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

Bayes limit

- p(S|x) = BD/(1 + BD) is what should be achieved to minimise cost, achieving classification with the fewest mistakes
- Fixing relative cost of background contamination and signal loss $q = C_B/(C_S + C_B)$, q = p(S|x) defines decision boundary:
 - signal-rich if $p(S|x) \ge q$
 - background-rich if p(S|x) < q
- Any function that approximates conditional class probability p(S|x) with negligible error reaches the Bayes limit



How to construct p(S|x)?

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

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

Now:

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

• Cutting on D(x) is equivalent to cutting on p(S|x), implying a corresponding (unknown) cut on p(S|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
 ⇒ generalisation ability







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Supervised learning

- Training events are labelled: N examples $(x, y)_1, (x, y)_2, \dots, (x, y)_N$ 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 ⇒ may discover unexpected features!

Google's research on building high-level features



A "giant" neural network

- At Google they trained a 9-layered NN with 1 billion connections
 - trained on 10 million 200×200 pixel images from YouTube videos
 - on 1000 machines (16000 cores) for 3 days, unsupervised learning
- Sounds big? The human brain has 100 billion (10¹¹) neurons and 100 trillion (10¹⁴) 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





Features extracted from such images



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

Machine learning



Finding the multivariate discriminant y = f(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:

$$R(w) = \frac{1}{N} \sum_{i=1}^{N} L(y_i, f(x_i, w))$$

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 $f(x, w_*)$, our estimate of y = f(x)

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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





































Quality of fit

- CPPM
- 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!

Quality of fit

- СРРМ
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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









Const. least squares fit, training RMSE = 0.915, test RMSE = 1.067





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





Quadr. least squares fit, training RMSE = 0.579, test RMSE = 0.723





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















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



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5) Summary
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|S) signal density
- b(x) = p(x|B) background density
- Cutting on D(x) is equivalent to cutting on probability p(S|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

- 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 *Kⁿ*: curse of dimensionality

Random grid search









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



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



Algorithm

- Better solutions more likely to be selected for mating and mutations, carrying their genetic code (cuts) from generation to generation
- Algorithm:
 - Create initial random population (cut ensemble)
 - 2 Select fittest individuals
 - 3 Create offsprings through crossover (mix best cuts)
 - Mutate randomly (change some cuts of some individuals)
 - Sepeat 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:

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

with vector of means μ and covariance matrix $\pmb{\Sigma}$

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

$$\ln B(x) = \lambda(x) \equiv \chi^2(\mu_B, \Sigma_B) - \chi^2(\mu_S, \Sigma_S)$$



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

- Fixed value of λ(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

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'Two moons' data



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Discriminant function with Gaussian fits

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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:

$$\lambda(x) = w \cdot x$$
 with $w \propto \Sigma^{-1}(\mu_S - \mu_B)$



- 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 ℝⁿ, but better separated if mapped to higher dimension space ℝ^H: h : x ∈ ℝⁿ → z ∈ ℝ^H
- Use hyper-planes to partition higher dim space: $f(x) = w \cdot h(x) + b$
- Example: $h: (x_1, y_2) \rightarrow (z_1, z_2, z_3) = (x_1^2, \sqrt{2}x_1x_2, x_2^2)$







СРРМ

Starting simple: separable data



- $w \cdot h(x) + b = 0$ (separating hyper-plane between red and blue)
- $w \cdot h(x_1) + b = +1$ (contains $h(x_1)$)
- $w \cdot h(x_2) + b = -1$ (contains $h(x_2)$)



- Subtract blue from red: $w \cdot (h(x_1) - h(x_2)) = 2$
- With unit vector $\hat{w} = w/||w||$: $\hat{w} \cdot (h(x_1) - h(x_2)) = 2/||w|| = m$
- 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: $w \cdot h(x) + b > 1$ for red dots $w \cdot h(x) + b < -1$ for blue dots

all correctly classified points will satisfy constraints:

$$y_i(w \cdot h(x_i) + b) \geq 1, \ \forall i = 1, \dots, N$$

• Using Lagrange multipliers $\alpha_i > 0$, cost function can be written:

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



Minimisation

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

$$C(\alpha) = \sum_{i=1}^{N} \alpha_i - \frac{1}{2} \sum_{i=1}^{N} \sum_{j=1}^{N} \alpha_i \alpha_j y_i y_j (h(x_i) \cdot h(x_j))$$

 At minimum of C(α), only non-zero α_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 $h(x_i) \cdot h(x_j)$
- Fortunately $h(x_i) \cdot h(x_j)$ are equivalent to some kernel function $K(x_i, x_j)$ that does the mapping and the scalar product:

$$C(\alpha) = \sum_{i=1}^{N} \alpha_i - \frac{1}{2} \sum_{i=1}^{N} \sum_{j=1}^{N} \alpha_i \alpha_j y_i y_j K(x_i, x_j)$$



Example



- In reality: do not know a priori the right kernel
- $\bullet\,\Rightarrow\,$ 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:

$$y_i(w \cdot h(x_i) + b) \geq 1 - \xi_i$$



with slack variables $\xi_i > 0$

- C(w, b, α, ξ) depends on ξ, modified C(α, ξ) 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 μ
- Density p(x) at point x approximated by:

$$p(x) pprox \hat{p}(x) = rac{1}{N} \sum_{j=1}^{N} K(x, \mu_j)$$





Choice of kernel

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

$$\mathcal{K}(x,\mu) = \prod_{i=1}^n \mathsf{Gaussian}(x_i|\mu,h_i)$$

each with bandwidth (width) h_i

Optimal bandwidth

- Too narrow: noisy approximation
- Too wide: loose fine structure
- In principle found by minimising risk function $R(\hat{p}, p) = \int (\hat{p}(x) p(x))^2 dx$
- For Gaussian densities:

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

Far from optimal for non-Gaussian densities





Example



Why does it work?

• When $N \to \infty$:

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

- $p(\mu)$: true density of x
- Kernel bandwidth getting smaller with N, so when $N \to \infty$, $K(x, \mu) \to \delta^n(x - \mu)$ 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



















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Discriminant function with Parzen fits, h = 0.01



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Discriminant function with Parzen fits, h = 0.25



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Training and test error rates



Neural networks





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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

Neural networks

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, λ is activation
- Neuron behaviour completely controlled by weights $w = \{w_0, \ldots, w_n\}$
- 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: $w^{(k+1)} = w^{(k)} \eta \frac{dE^{(k)}}{dw}$
- $\frac{\partial E}{\partial w_j} = \sum_{n=1}^{N} -(t^{(n)} y^{(n)}) x_j^{(n)}$ 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 $C(I_n)$ denote the space of continuous functions on the n-dimensional hypercube. Then, for any given function $f \in C(I_n)$ and $\varepsilon > 0$ there exists an integer M and sets of real constants w_j, w_{ij} where i = 1, ..., n and j = 1, ..., M such that

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

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

$$y_k(x,w) = \sum_{j=0}^{M} w_{kj}^{(2)} \underbrace{\sigma\left(\sum_{i=0}^{n} w_{ji}^{(1)} x_i\right)}_{z_i}$$







Can fit any function: examples

- 1 input (training data), 1 output
- 3 hidden neurons on one hidden layer



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Backpropagation

- Training means minimising error function *E*(*w*)
- For single neuron: $\frac{dE}{dw_k} = (y t)x_k$
- One can show that for a network:

$$\frac{dE}{dw_{ji}} = \delta_j z_i, \text{ where }$$

$$\delta_k = (y_k - t_k)$$
 for output neurons
 $\delta_j \propto \sum_k w_{kj} \delta_k$ otherwise

 z_i

• As before, weights can be regularised:

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

• Hence errors are propagated backwards



Regularisation





• 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 + rand(-0.05, +0.05)
- Of course doesn't work as well if noise is not symmetric



Tricks of the trade

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- 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, ..., (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|T) to it
- If p(w₁|T) > p(w₂|T), then associated function f(x, w₁) more compatible with training data T than function f(x, w₂)
- Posterior density p(w|T) is final result of Bayesian inference
- BNN is the predictive distribution

$$p(y|x,T) = \int p(y|x,w)p(w|T)dw$$

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:

$$y(x) = \int zp(z|x, T)dz$$
$$= \int f(x, w)p(w|T)dw$$

• Why? For classification $p(y|x, w) = f(x, w)^{y} (1 - f(x, w))^{1-y}$

• for
$$y = 1$$
: $p(y|x, w) = f(x, w)$

• for
$$y = 0$$
: $p(y|x, w) = 1 - f(x, w)$

• so only f(x, w) contributes to the mean

• Example usage:

$$f(x, w) = \frac{1}{1 + e^{-g(x,w)}}$$

$$g(x, w) = b + \sum_{j=1}^{H} v_j \tanh(a_j + \sum_{i=1}^{n} u_{ij}x_i)$$

with H hidden nodes



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|T)
- $y(x) \approx \frac{1}{M} \sum_{m=1}^{M} f(x, w_m)$
- 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 $f(x, w_k)$
- thick curve: average, which approximates D(x)



Details tomorrow

Summary of MVA techniques



Criteria		Classifiers								
		Cuts	Likeli- hood	PDERS / k-NN	H-Matri x	Fisher	MLP	BDT	RuleFit	SVM
Perfor- mance	no / linear correlations	•	C	0	(0	0	•	0	0
	nonlinear correlations	(8	\odot	8	8	\odot	\odot	(0
Speed	Training	8	C	\odot	\odot	\odot	(8	(8
	Response	\odot	\odot	⊗/☺	\odot	\odot	0	•	(:
Robust -ness	Overtraining	\odot	((\odot	\odot	8	8	((
	Weak input variables	\odot	\odot	8	\odot	\odot	(((3
Curse of dimensionality		8	\odot	8	\odot	\odot	(\odot	(:
Transparency		\odot	\odot	•	\odot	\odot	8	8	8	8

(according to TMVA authors)

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• 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://www.phi-t.de/mousegame/index_eng.html



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