PCML Cheat Sheet

1 Math Prerequisites

- Bayes rule

$$p(A,B) = \underbrace{p(A|B)}_{\text{Lik.}} \underbrace{p(B)}_{\text{Prior}} = \underbrace{p(B|A)}_{\text{Post}} \underbrace{p(A)}_{\text{Marg. Lik.}}$$

- Gaussian distribution

$$\mathcal{N}(x|\mu,\sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp(-\frac{(x-\mu)^2}{2\sigma^2})$$

$$\mathcal{N}(\mathbf{x}|\boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{1}{\sqrt{2\pi|\boldsymbol{\Sigma}|}} \exp(-\frac{1}{2}(\mathbf{x} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1}(\mathbf{x} - \boldsymbol{\mu}))$$

Production of independent variables:

$$\operatorname{Var}(XY) = \mathbb{E}(X^2) \,\mathbb{E}(Y^2) - [\mathbb{E}(X)]^2 [\mathbb{E}(Y)]^2$$

- Covariance matrix of a data vector x

$$\Sigma = \frac{1}{N} \sum_{n=1}^{N} (\mathbf{x}_n - \mathbb{E}(\mathbf{x})) (\mathbf{x}_n - \mathbb{E}(\mathbf{x}))^T$$

- A function f(x) is convex, if for any $x_1, x_2 \in \mathbf{X}$ and for any $0 \le \lambda \le 1$, we have :

$$f(\lambda x_1 + (1 - \lambda)x_2) \le \lambda f(x_1) + (1 - \lambda)f(x_2)$$

- Strictly convex if the inequality is strict.
- Sums of convex functions are also convex.
- A function with the form log-sum-exp is convex.
- The Hessian is related to the convexity: a twice differentiable function is convex i-o-if the Hessian is positive definite.
- The Hessian of a convex function is positive semi-definite and for a strictly-convex function it is positive definite.

$$\mathbf{H}_{i,j} = d^2 f / dx_i dx_j$$

1.2 Linear Algebra

- Column $\mathbf{x} \in \mathbb{R}^n$, rows \mathbf{x}^T , matrix $\mathbf{A} \in \mathbb{R}^{m \times n}$
- $\mathbf{x}^T \mathbf{x}$ is a scalar, $\mathbf{x} \mathbf{x}^T$ is a matrix
- \mathbf{A}^{-1} exist if \mathbf{A} is full rank $(\mathbf{A}\mathbf{B})^T = \mathbf{B}^T \mathbf{A}^T$
- Condition number of a function measures how much the output value can change for a small change in the input. A matrix with a high condition number is said to be **ill-conditioned**. If **A** is normal $(A^T A = AA^T)$ then

$$k(\mathbf{A}) = \left| rac{\lambda_{max}(\mathbf{A})}{\lambda_{min}(\mathbf{A})}
ight|$$
– A positive definite matrix is **symmetric** with

- all positive eigenvalues
- The real symmetric $N \times N$ matrix \mathbf{V} is said to be positive semidefinite if

$$\mathbf{a}^T \mathbf{V} \mathbf{a} > 0$$

- for any real $N \times 1$ vector a.
- positive definite if $\mathbf{a}^T \mathbf{V} \mathbf{a} > 0$
- Cost of matrix inversion: $O(n^3) \rightarrow O(n^{2.372})$
- det(A) using LU decomposition: O(n³)

2 Cost functions

- Cost functions are used to learn parameters that explain the data well.
- It is essential to make sure that a global minimum exist \rightarrow lower bounded

Mean square error (MSE):

$$MSE(\boldsymbol{w}) = \sum_{n=1}^{N} (y_n - f(\mathbf{x}_n))^2$$

- MSE is convex thus it has only one global
- MSE is not good when outliers are present.

Mean Absolute Error (MAE):

$$MAE = \sum_{n=1}^{N} |y_n - f(\mathbf{x}_n)|$$

$$Huber = \left\{ \begin{array}{cc} \frac{1}{2}z^2 & , |z| \leq \delta \\ \delta |z| - \frac{1}{2}\delta^2 & , |z| > \delta \end{array} \right.$$

Huber loss is convex, differentiable, and also

Tukey's bisquare loss

$$L(z) = \begin{cases} z(\delta^2 - z^2)^2 &, |z| < \delta \\ 0 &, |z| \ge \delta \end{cases}$$

Non-convex, non-diff., but robust to outliers. Hinge loss

 $Hinge = [1 - y_n f(\mathbf{x}_n)]_+ = \max(0, 1 - y_n f(\mathbf{x}_n))$ Logistic loss

$$Logistic = log(1 - exp(y_n f(\mathbf{x}_n)))$$

3 Regression

- **Data** consists of N pairs (y_n, \mathbf{x}_n)
- 2. \mathbf{x}_n is a vector of D inputs
- Prediction: predict the ouput for a new input
- Interpretation: understand the effect of inputs
- Outliers are data that are far away from most of the other examples.

3.1 Linear Regression

- Model that assume linear relationship between inputs and the ouput.

$$y_n \approx f(\mathbf{x}_n) := w_0 + w_1 x_{n1} + \dots = \omega_0 + \mathbf{x}_n^T \mathbf{w}$$
 with \mathbf{w} the parameters of the model.

Variance grows only linearly with dimensionality

4 Optimization

4.1 Grid search

- Compute the cost over a grid of M points to find the minimum
- Exponential Complexity
- Hard to find a good range of values

4.2 Gradient Descent

- Gradient descent uses only first-order information and takes steps in the direction of the gradient
- Given a cost function $\mathcal{L}(\boldsymbol{w})$ we wish to find \boldsymbol{w} that minimizes the cost:

$$\min_{\boldsymbol{x}} \mathcal{L}(\boldsymbol{w})$$

4.3 Batch Gradient Descent

- Take steps in the opposite direction of the gradient

$$\boldsymbol{w}^{(t+1)} \leftarrow \boldsymbol{w}^{(t)} - \gamma \nabla \mathcal{L}(\boldsymbol{w}^{(t)})$$

with $\gamma > 0$ the learning rate. With γ too big, method might diverge. With γ too small, convergence is slow.

4.4 Gradients for MSE

We define the error vector e:

$$e := y - Xw$$

- and MSE as follows:

$$\mathcal{L}(\boldsymbol{w}) = \frac{1}{2N} \sum_{n=1}^{N} (y_n - \tilde{\mathbf{x}}_n^T \boldsymbol{w})^2 = \frac{1}{2N} \mathbf{e}^T \mathbf{e}$$

- then the gradient is given by

$$\nabla \mathcal{L}(w) = -\frac{1}{N} \mathbf{X}^T \mathbf{e}$$

- Optimality conditions

- 1. necessary: gradient equal zero: $\frac{d\mathcal{L}(\boldsymbol{w}^*)}{d\boldsymbol{w}} = 0$ 2. sufficient: Hessian matrix is positive definite:

$$\mathbf{H}(\mathbf{w}^*) = \frac{d^2 \mathcal{L}(\mathbf{w}^*)}{T}$$

- $\mathbf{H}(\boldsymbol{w}^*) = \frac{d^2 \mathcal{L}(\boldsymbol{w}^*)}{d \boldsymbol{w} d \boldsymbol{w}^T}$ Very sensitive to illconditioning. Therefore, always normalize your feature otherwise step-size selection is difficult since different directions might move at different speed.
- Complexity: O(NDI) with I the number of

4.5 Stochastic Gradient Descent

In ML, most cost functions are formulated as a sum over the training examples:

$$\mathcal{L}(\boldsymbol{w}) = \frac{1}{N} \sum_{n=1}^{N} \mathcal{L}_n(\boldsymbol{w})$$

⇒ SGD algo is given by update rule:

$$\boldsymbol{w}^{(t+1)} \leftarrow \boldsymbol{w}^{(t)} - \gamma \nabla \mathcal{L}_n(\boldsymbol{w}^{(t)})$$

Idea: Cheap but unbiased estimate of grad

$$\mathbb{E}[\boldsymbol{\nabla}\mathcal{L}_n(\boldsymbol{w})] = \boldsymbol{\nabla}(\boldsymbol{w})$$

4.6 Mini-batch SGD

Update direction $(B \subseteq [N])$:

$$g^{(t)} := \frac{1}{|B|} \sum_{n \in B} \nabla \mathcal{L}_n(w^{(t)})$$

Update rule: $\mathbf{w}^{(t+1)} \leftarrow \mathbf{w}^{(t)} - \gamma \mathbf{g}^{(t)}$

4.7 Subgradients (Non-Smooth OPT)

A vector $\mathbf{g} \in \mathbb{R}^D$ s.t.

$$\mathcal{L}(u) \geq \mathcal{L}(w) + g^T(u - w) \quad \forall u$$

is the subgradient to $\mathcal L$ at $\boldsymbol w$. If $\mathcal L$ is differentiable

5 Least Squares

- In some cases, we can compute the minimum of the cost function analytically.
- use the first optimality conditions:

$$\nabla L(\boldsymbol{w}^*) = 0 \Rightarrow \boldsymbol{X}^T \boldsymbol{e} = \boldsymbol{X}^T (\boldsymbol{y} - \boldsymbol{X} \boldsymbol{w}) = 0$$

– When $\mathbf{X}^T\mathbf{X}$ is invertible, we have the closed-form expression

$$\mathbf{w}^* = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$$

thus we can predict values for a new \mathbf{x}_m $y_m := \mathbf{x}_{\mathbf{m}}^{\mathbf{T}} \mathbf{w}^* = \mathbf{x}_{\mathbf{m}}^{\mathbf{T}} (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$

$$ym \leftarrow \mathbf{A}_{\mathbf{m}} \mathbf{w} - \mathbf{A}_{\mathbf{m}} (\mathbf{A}, \mathbf{A})$$
The **Gram matrix** $\mathbf{X}^T \mathbf{X}$ is positive definite and is also invertible iff \mathbf{X} has full column rank. Complexity: $O(ND^2 + D^3) \equiv O(ND^2)$

X can be rank deficient when D > N or when the comlumns $\bar{\mathbf{x}}_d$ are nearly collinear. In this case, the matrix is ill-conditioned, leading to

6 Maximum Likelihood

Let define our mistakes ε_n ~ N(0, σ²).

$$\rightarrow y_n = \mathbf{x}_n^T \mathbf{w} + \epsilon_n$$

Another way of expressing this:

$$\begin{aligned} p(\mathbf{y}|\mathbf{X}, \mathbf{w}) &= \prod_{n=1}^{N} p(y_n|\mathbf{x}_n, \mathbf{w}) \\ &= \prod_{n=1}^{N} \mathcal{N}(y_n|\mathbf{x}_n^T \mathbf{w}, \sigma^2) \end{aligned}$$

which defines the likelihood of observating y given ${\bf X}$ and ${\bf w}$

Define cost with log-likelihood

$$\begin{split} \mathcal{L}_{lik}(\boldsymbol{w}) &= \log p(\mathbf{y}|\mathbf{X}, \boldsymbol{w}) \\ &= -\frac{1}{2\sigma^2} \sum_{i}^{N} (y_n - \mathbf{x}_n^T \boldsymbol{w})^2 + cnst \end{split}$$

 Maximum likelihood estimator (MLE) gives another way to design cost functions

her way to design cost functions
$$\operatorname{argmin} \mathcal{L}_{MSE}(w) = \operatorname{argmax} \mathcal{L}_{lik}(w)$$

- MLE can also be interpreted as finding the model under which the observed data is most likely to have been generated from.

$$p(y_n|\mathbf{x}_n, \mathbf{w}) = \frac{1}{2} e^{-\frac{1}{b}|y_n - \mathbf{x}_n^T \mathbf{w}|}$$

$$\sum_{n} \log p(y_n | \mathbf{x}_n, \mathbf{w}) = \sum_{n} |y_n - \mathbf{x}_n^T \mathbf{w}| + cnst$$

- Linear models usually underfit. One way is to use nonlinear basis functions instead.

$$y_n = w_0 + \sum_{j=1}^{M} w_j \phi_j(\mathbf{x}_n) = \tilde{\phi}(\mathbf{x}_n)^T \mathbf{w}$$

This model is linear in \boldsymbol{w} but nonlinear in \boldsymbol{x} . Note that the dimensionality is now M, not D. Polynomial basis

$$\phi(x_n) = [1, x_n, x_n^2, ..., x_n^M]$$

- The least square solution becomes $\mathbf{w}_{lse}^* = (\mathbf{\tilde{\Phi}}^T \mathbf{\tilde{\Phi}})^{-1} \mathbf{\tilde{\Phi}}^T \mathbf{y}$

Complex models overfit easily. Thus we can choose simpler models by adding a regularization term which penalizes complex

$$\min_{\boldsymbol{w}} \left(\mathcal{L}(\boldsymbol{w}) + \frac{\lambda}{2N} \sum_{j=1}^{M} w_j^2 \right)$$

$$\boldsymbol{w}^* = \operatorname*{argmin}_{\boldsymbol{w}} \left(\frac{1}{2} (\mathbf{y} - \mathbf{X} \boldsymbol{w})^T (\mathbf{y} - \mathbf{X} \boldsymbol{w}) + \frac{\lambda}{2} \boldsymbol{w}^T \boldsymbol{w} \right)$$

Note that w₀ is not penalized.
By differentiating and setting to zero we get

$$egin{align} w_{ridge} &= (ilde{f \Phi}^T ilde{f \Phi} + {f \Lambda})^{-1} ilde{f \Phi}^T {f y} \ f \Lambda &= \left[egin{array}{cc} 0 & 0 \ 0 & \lambda ilde{I}_m \end{array}
ight] \end{array}$$

Ridge regression improves the condition number of the Gram matrix since the eigenvalues of $(\mathbf{\tilde{\Phi}}^T\mathbf{\tilde{\Phi}} + \lambda I_m)$ are at least λ

Maximum-a-posteriori (MAP) estimator:

- Maximizes the product of the likelihood and the prior.

$$\boldsymbol{w}_{MAP} = \operatorname*{argmax} \left(p(\mathbf{y}|\mathbf{X}, \boldsymbol{\Lambda}) p(\boldsymbol{w}|\boldsymbol{\Sigma}) \right)$$

- Assume $w_0 = 0$

$$oldsymbol{w}_{ridge} = \operatorname*{argmax}_{oldsymbol{w}} \left(\log \left[\prod_{n=1}^{N} \mathcal{N}(y_n | \mathbf{x}_n^T w, \mathbf{\Lambda}) \right] \right)$$

Lasso regularizer forces some w_i to be strictly

$$\begin{split} \min_{\pmb{w}} \ \frac{1}{2N} \sum_{n=1}^{N} (y_n - \tilde{\pmb{\phi}}(\mathbf{x}_n)^T \pmb{w})^2, \\ \text{such that } \sum_{n=1}^{M} |w_i| \leq \tau \end{split}$$

8 Cross-Validation

We should choose λ to minimize the mistakes that will be made in the future.

- We split the data into train and validation sets and we pretend that the validation set is the future data. We fit our model on the training set and compute a prediction-error on the validation set. This gives us an estimate of the generalization error.
- K-fold cross validation randomly partition the data into K groups. We train on K-1groups and test on the remaining group. We repeat this until we have tested on all K sets. We then average the results.
- Cross-validation returns an unbiased estimate of the generalization error and its variance.

9 Bias-Variance decomposition

around their average

- The expected test error can be expressed as the sum of two terms

- Squared bias: The average shift of the

predictions - Variance: measure how data points vary

expected loss =
$$(bias)^2$$
 + variance + noise

- Both model bias and estimation bias are
- important Ridge regression increases estimation bias while
- reducing variance Increasing model complexity increases test error
- Small λ → low bias but large variance
 Large λ → large bias but low variance

Large
$$\lambda \to \text{large bias but low variance}$$

$$err = \sigma^2 + E[f_{lse} - E[f_{lse}]]^2 + [f_{true} - E[f_{lse}]]^2$$

10 Logistic Regression

- Classification relates input variables x to
- discrete output variable y Binary classifier: we use y = 0 for C_1 and
 - y = 1 for C_2 .

Can use least-squares to predict
$$\hat{y}_*$$

$$\hat{y} = \begin{cases} \mathbf{C}_1 & \hat{y}_* < 0.5 \\ \mathbf{C}_2 & \hat{y}_* \ge 0.5 \end{cases}$$

- Logistic function

- The probabilistic model:

$$\sigma(x) = \frac{\exp(x)}{1 + \exp(x)}$$

$$p(y_n = \mathbf{C}_1 | \mathbf{x}_n) = \sigma(\mathbf{x}^T \mathbf{w})$$
$$p(y_n = \mathbf{C}_2 | \mathbf{x}_n) = 1 - \sigma(\mathbf{x}^T \mathbf{w})$$

$$p(\mathbf{y}|\mathbf{X}, \boldsymbol{w}) = \prod_{n=1}^{N} \sigma(\boldsymbol{x}_{n}^{T} \boldsymbol{w})^{y_{n}} (1 - \sigma(\boldsymbol{x}_{n}^{T} \boldsymbol{w}))^{1 - y_{n}}$$

- The log-likelihood:

$$\mathcal{L}_{MLE}(\boldsymbol{w}) = \sum_{n=1}^{N} \left(y_n \boldsymbol{x}_n^T \boldsymbol{w} - \log(1 + \exp(\boldsymbol{x}_n^T \boldsymbol{w})) \right)$$

- We can use the fact that

$$\frac{d}{dx}\log(1+\exp(x)) = \sigma(x)$$

$$\mathbf{g} = \frac{d\mathcal{L}}{d\mathbf{w}} = \sum_{n=1}^{N} \left(\mathbf{x}_n y_n - \mathbf{x}_n \sigma(\mathbf{x}_n^T \mathbf{w}) \right)$$
$$= -\mathbf{X}^T [\sigma(\mathbf{X} \mathbf{w}) - \mathbf{y}]$$

- The negative of the log-likelihood $-\mathcal{L}_{mle}(\boldsymbol{w})$ is

Hessian of the log-likelihood

- We know that

$$\frac{d\sigma(t)}{t} = \sigma(t)(1 - \sigma(t))$$

 $w_{ridge} = \underset{\boldsymbol{w}}{\operatorname{argmax}} \left(\log \left[\prod_{n=1}^{N} \mathcal{N}(y_n | \mathbf{x}_n^T \boldsymbol{w}, \boldsymbol{\Lambda}) \right. \right) \\ \left. \begin{array}{l} \frac{d\sigma(t)}{dt} = \sigma(t) (1 - \sigma(t)) \\ \mathcal{N}(\boldsymbol{w} | \boldsymbol{0}, \boldsymbol{H}) \end{array} \right) \right) \\ \text{So regularizer forces are simple solutions}$ $\mathbf{H}(\boldsymbol{w}) = -\frac{d\mathbf{g}(\boldsymbol{w})}{d\boldsymbol{w}^T} = \sum_{n=1}^{N} \frac{d}{d\boldsymbol{w}^T} \sigma(\boldsymbol{x}_n^T \boldsymbol{w}) \boldsymbol{x}_n$

$$n=1 \text{ or } n=1$$

$$= \sum_{n=1}^{N} \mathbf{x}_n \sigma(\mathbf{x}_n^T \mathbf{w}) (1 - \sigma(\mathbf{x}_n^T \mathbf{w})) \mathbf{x}_n^T$$

where **S** is a $N \times N$ diagonal matrix with

$$S_{nn} = \sigma(\mathbf{x}_n^T \mathbf{w})(1 - \sigma(\mathbf{x}_n^T \mathbf{w}))$$

 The negative of the log-likelihood is not strictly convey Newton's Method Uses second-order information and takes

steps in the direction that minimizes a quadratic approximation
$$\mathcal{L}(\boldsymbol{w}) = \mathcal{L}(\boldsymbol{w}^{(k)}) + \mathbf{g}_k^T(\boldsymbol{w} - \boldsymbol{w}^{(k)})$$

 $+(\boldsymbol{w}-\boldsymbol{w}^{(k)})^T\mathbf{H}_{k}(\boldsymbol{w}-\boldsymbol{w}^{(k)})$

 $\boldsymbol{w}^{k+1} = \boldsymbol{w}^{(k)} - \alpha_k \boldsymbol{H}_{l}^{-1} \boldsymbol{g}_{l_k}$ where \mathbf{g}_k is the gradient and α_k the

learning rate.

$$\begin{aligned} & - & \text{Complexity: } O((ND^2 + D^3)I) \\ & \textbf{Penalized Logistic Regression} \\ & \min_{\boldsymbol{w}} \left(- \sum_{n=1}^{N} \log p(y_n | \mathbf{x}_n^T \boldsymbol{w}) + \lambda \sum_{n=1}^{D} w_d^2 \right) \end{aligned}$$

11 Generalized Linear Model

Exponential family distribution
$$p(\mathbf{y}|\boldsymbol{\eta}) = \frac{h(y)}{\sigma} \exp(\boldsymbol{\eta}^T \boldsymbol{\phi}(\mathbf{y}) - A(\boldsymbol{\eta}))$$

- Bernoulli distribution
$$p(y|\mu) = \mu^{y}(1-\mu)^{1-y}$$

$$= \exp(y \log(\frac{\mu}{1-\mu} + \log(1-\mu)))$$

- there is a relationship between η and μ throught the link function

ght the link function
$$\eta = \log(\frac{\mu}{1 - \mu}) \leftrightarrow \mu = \frac{e^{\eta}}{1 + e^{\eta}}$$

 Note that μ is the mean parameter of y Relationship between the mean μ and η is defined using a link function q

$$\begin{split} \eta &= \mathbf{g}(\boldsymbol{\mu}) \Leftrightarrow \boldsymbol{\mu} = \mathbf{g}^{-1}(\boldsymbol{\eta}) \\ \text{First and second derivatives of } A(\boldsymbol{\eta}) \text{ are related to the mean and the variance} \\ \frac{dA(\boldsymbol{\eta})}{d\boldsymbol{\eta}} &= \mathbb{E}[\phi(\boldsymbol{\eta})], \quad \frac{d^2A(\boldsymbol{\eta})}{d\boldsymbol{\eta}^2} = \mathrm{Var}[\phi(\boldsymbol{\eta})] \end{split}$$

- $A(\eta)$ is convex - The generalized maximum likelihood cost to

$$\min_{\boldsymbol{w}} \mathcal{L}(\boldsymbol{w}) = -\sum_{n=1}^{N} \log(p(y_n | \boldsymbol{x}_n^T \boldsymbol{w}))$$

where $p(y_n|\boldsymbol{x}_n^T\boldsymbol{w})$ is an exponential family

- We obtain the solution

$$\frac{d\mathcal{L}}{d\mathbf{w}} = \mathbf{X}^T [\mathbf{g}^{-1}(\boldsymbol{\eta}) - \boldsymbol{\phi}(\mathbf{y})]$$

12 k-Nearest Neighbor (k-NN)

- The k-NN prediction for x is

$$f(\mathbf{x}) = \frac{1}{k} \sum_{\mathbf{x}_n \in nbh_k(\mathbf{x})} y_n$$

where $nbh_k(\mathbf{x})$ is the neightborhood of \mathbf{x} defined by the k closest points \mathbf{x}_n in the training data

- Curse of dimensionality: Generalizing correctly becomes exponentially harder as the dimensionality grows.
- Gathering more inputs variables may be a bad

13 Support Vector Machine

- Combination of the kernel trick plus a modified loss function (Hinge loss)
- Solution to the dual problem is sparse and non-zero entries will be our support vectors.
- Kernelised feature vector where μ_k are centroids

$$\phi(\mathbf{x}) = [k(\mathbf{x}, \boldsymbol{\mu}_1), ..., k(\mathbf{x}, \boldsymbol{\mu}_K)]$$

- In practice we'll take a subset of data points to be prototype -> sparse vector machine.
- Assume $y_n \in \{-1, 1\}$
- SVM optimizes the following cost

$$\mathcal{L}(\boldsymbol{w}) = \min_{\boldsymbol{w}} \sum_{n=1}^{N} [1 - y_n \tilde{\boldsymbol{\phi}}_n^T \boldsymbol{w}]_+ + \frac{\lambda}{2} \sum_{j=1}^{M} w_j^2$$

- Minimum doesn't change with a rescaling of w choose the hyperplane so that the distance from it to the nearest data point on each side is maximized
- Duality
 - Hard to minimize g(w) so we define

$$\mathcal{L}(\boldsymbol{w}) = \max_{\boldsymbol{\alpha}} G(\boldsymbol{w}, \boldsymbol{\alpha})$$

- we use the property that

$$C[v_n]_+ = \max(0, Cv_n) = \max_{\alpha_n \in [0, C]} \alpha_n v_n$$

- We can rewrite the problem as

$$\min_{\boldsymbol{w}} \max_{\boldsymbol{\alpha} \in [0,C]^N} \sum_{n=1}^N \alpha_n (1 - y_n \boldsymbol{\phi}_n^T \boldsymbol{w}) + \frac{1}{2} \sum_{j=1}^M w_j^2$$

- This is differentiable, convex in w and
- concave in α

$$\min_{\boldsymbol{w}} \max_{\boldsymbol{\alpha}} G(\boldsymbol{w}, \boldsymbol{\alpha}) = \max_{\boldsymbol{\alpha}} \min_{\boldsymbol{w}} G(\boldsymbol{w}, \boldsymbol{\alpha})$$

because G is convex in w and concave in α .

– Derivative w.r.t.
$$m{w}$$
:
$$\nabla_{m{w}} G(m{w}, \pmb{\alpha}) = -\sum_{n=1}^{N} \alpha_n y_n \pmb{x}_n + \lambda \pmb{w}$$

- Equating this to 0, we get:

$$w(\alpha) = \frac{1}{\lambda} \sum_{n=1}^{N} \alpha_n y_n x_n = \frac{1}{\lambda} XY\alpha$$

Plugging w* back in the dual problem

$$\max_{\boldsymbol{\alpha} \in [0,1]^N} \boldsymbol{\alpha}^T \mathbf{1} - \frac{1}{2\lambda} \boldsymbol{\alpha}^T \mathbf{Y} \boldsymbol{X}^T \boldsymbol{X} \mathbf{Y} \boldsymbol{\alpha}$$

- This is a differentiable least-squares problem. Optimization is easy using Sequential Minimal Optimization. It is also naturally kernelized with $\mathbf{K} = \mathbf{X}^T \mathbf{X}$
- The solution α is sparse and is non-zero only for the training examples that are instrumental in determining the decision boundary.

14 Kernel Ridge Regression

- The following is true for ridge regression

$$\mathbf{w}^* = (\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I}_D)^{-1} \mathbf{X}^T \mathbf{y}$$

$$= \mathbf{X}^T (\mathbf{X} \mathbf{X}^T + \lambda \mathbf{I}_N)^{-1} \mathbf{y} = \mathbf{X}^T \boldsymbol{\alpha}^*$$
(1)

- Complexity of computing w: (1)
- $O(D^2N + D^3)$, (2) $O(DN^2 + N^3)$
- Thus we have

$$w^* = \mathbf{X} \boldsymbol{\alpha}^*, \quad \text{with } w^* \in \mathbb{R}^D \text{ and } \boldsymbol{\alpha}^* \in \mathbb{R}^N$$

The representer theorem allows us to write an equivalent optimization problem in terms of α .

$$\boldsymbol{\alpha} = \operatorname*{argmax}_{\boldsymbol{\alpha}} \left(-\frac{1}{2} \boldsymbol{\alpha}^T (\boldsymbol{\mathbf{X}}^T \boldsymbol{\mathbf{X}} + \lambda \boldsymbol{\mathbf{I}}_N) \boldsymbol{\alpha} + \boldsymbol{\alpha}^T \boldsymbol{\mathbf{y}} \right)$$

- $\mathbf{K} = \mathbf{X}\mathbf{X}^T$ is called the **kernel matrix** or Gram matrix.
- If K is positive definite, then it's called a Mercer Kernel
- $\begin{array}{ll} & \mathbf{K}_{i,j} = k(\mathbf{x}_i, \mathbf{x}_j) \\ & \text{If the kernel is Mercer, then there exists a} \end{array}$ function $\phi(\mathbf{x})$ s.t.

$$k(\mathbf{x}, \mathbf{x}') = \phi(\mathbf{x})^T \phi(\mathbf{x}')$$

- Kernel trick:
- We can work directly with K and never have to worry about X
- Replace (x, x') with k(x, x').
- Kernel function can be interpreted as a measure of similarity
- The evaluation of a kernel is usually faster with k than with d
- Kernelized rigde regression might be computationally more efficient in some cases
- Radial Basis function kernel (RBF)

$$k(\mathbf{x}, \mathbf{x'}) = \exp(-\frac{1}{2}(\mathbf{x} - \mathbf{x'})^T(\mathbf{x} - \mathbf{x'}))$$

- Properties of a kernel to ensure the existance of a corresponding d:
- **K** should be symmetric: $k(\mathbf{x}, \mathbf{x}') = k(\mathbf{x}', \mathbf{x})$ - K should be positive semidefinite.

$$\mathbf{y} = \mathbf{w}^T \mathbf{x} = \sum_{i=1}^K \alpha_i \mathbf{x}_i^T \mathbf{x} = \sum_{i=1}^K \alpha_i k(\mathbf{x}, \mathbf{x}_i)$$

15 K-means

- Unsupervised learning: Represent particular input patterns in a way that reflects the statistical structure of the overall collections of input partterns.
- Cluster are groups of points whose inter-point distances are small compared to the distances

$$\min_{\mathbf{r}, \boldsymbol{\mu}} \mathcal{L}(\mathbf{r}, \boldsymbol{\mu}) = \sum_{k=1}^K \sum_{n=1}^N r_{nk} ||\mathbf{x}_n - \boldsymbol{\mu}_k||_2^2$$

- such that $r_{nk} \in \{0, 1\}$ and $\sum_{k=1}^{K} r_{nk} = 1$ K-means algorithm:
- Initialize μ_k , then iterate
- 1. For all n, compute \mathbf{r}_n given $\boldsymbol{\mu}$

$$r_{nk} = \begin{cases} & \text{if } k = \operatorname{argmin}_{j} ||\mathbf{x}_{n} - \boldsymbol{\mu}||_{2}^{2} \\ & \text{otherwise} \end{cases}$$

$$\mu_k = \frac{\sum_{n=1}^{N} r_{nk} \mathbf{x}_n}{\sum_{n=1}^{N} r_{nk}}$$

- A good initialization procedure is to choose the prototypes to be equal to a random subset of Kdata points.
- Probabilistic model

$$p(\mathbf{r}, \boldsymbol{\mu}) = \prod_{n=1}^{N} \prod_{k=1}^{K} \left[\mathcal{N}(\mathbf{x}_{n} | \boldsymbol{\mu}_{k}, \mathbf{I}) \right]^{r_{n} k}$$

Computation can be heavy, each example can belong to only on cluster and clusters have to be spherical.

16 Gaussian Mixture Models

- Clusters can be spherical using a full covariance matrix instead of isotropic covariance.

$$p(\mathbf{X}|\boldsymbol{\mu},\boldsymbol{\Sigma},\mathbf{r}) = \prod_{n=1}^{N} \prod_{k=1}^{K} \left[\mathcal{N}(\mathbf{x}_{n}|\boldsymbol{\mu}_{k},\boldsymbol{\Sigma}_{k}) \right]^{r} nk$$

Soft-clustering: Points can belong to several cluster by defining r_n to be a random variable

$$p(r_{nk} = 1) = \pi_k \text{ where } \pi_k > 0, \forall k, \sum_{k=1}^K \pi_k = 1$$

Joint distribution of Gaussian mixture model

$$\begin{split} p(\mathbf{X}, \mathbf{r} | \boldsymbol{\mu}, \boldsymbol{\Sigma}, \boldsymbol{\pi}) &= \prod_{n=1}^{N} \left[p(\mathbf{x}_n | \mathbf{r}_n, \boldsymbol{\mu}, \boldsymbol{\Sigma}) p(\mathbf{r}_n | \boldsymbol{\pi}) \right] \\ &= \left[\prod_{k=1}^{K} \left[\left(\mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right)^r nk \right] \prod_{k=1}^{K} \left[r_k (\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right] \right] \end{split}$$

- r_n are called *latent* unobserved variables Unknown parameters are given by $\theta = \{\mu, \Sigma, \pi\}$
- We get the marginal likelihood by marginalizing r_n out from the likelihood

$$\begin{split} p(\mathbf{x}_n|\boldsymbol{\theta}) &= \sum_{k=1}^K p(\mathbf{x}_n, r_{nk} = 1|\boldsymbol{\theta}) \\ &= \sum_{k=1}^K p(r_{nk} = 1|\boldsymbol{\theta}) p(\mathbf{x}_n|r_{nk} = 1, \boldsymbol{\theta}) \\ &= \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}_n|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \end{split}$$

- Without a latent variable model, number of parameters grow at rate O(N)
- After marginalization, the growth is reduced to $O(D^2K)$
- To get maximum likelihood estimate of θ , we

$$\max_{\pmb{\theta}} \sum_{n=1}^{N} \log \sum_{k=1}^{K} \pi_k \mathcal{N}(\mathbf{x}_n | \pmb{\mu}_k, \pmb{\Sigma}_k)$$

17 Expectation Maximization Algorithm

- [ALGORITHM] Start with $\theta^{(1)}$ and iterate
- 1. Expectation step: Compute a lower bound to the cost such that it is tight at the previous $\boldsymbol{\theta}^{(i)}$ with equality when,

$$\gamma(r_{nk}) = \frac{\pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}$$

2. Maximization step: Update θ

$$\boldsymbol{\theta}^{(i+1)} = \operatorname*{argmax}_{\boldsymbol{\theta}} \mathcal{L}(\boldsymbol{\theta}, \boldsymbol{\theta}^{(i)})$$

$$\mu_k^{(i+1)} = \frac{\sum_{n=1}^{N} \gamma^{(i)}(r_{nk}) \mathbf{x}_n}{\sum_{n=1}^{N} \gamma^{(i)}(r_{nk})}$$

$$\mathbf{\Sigma}_{k}^{(i+1)} = \frac{\sum_{n=1}^{N} \gamma^{(i)}(r_{nk})(\mathbf{x}_{n} - \boldsymbol{\mu}_{k}^{(i+1)})(\mathbf{x}_{n})}{\sum_{n=1}^{N} \gamma^{(i)}(r_{nk})}$$

$$\pi_k^{(i+1)} = \frac{1}{N} \sum_{n=1}^{N} \gamma^{(i)}(r_{nk})$$

18 Matrix factorization

- We have D movies and N users
- $\mathbf X$ is a matrix $D\times N$ with x_{dn} the rating of n'th user for d'th movie.
- We project data vectors x_n to a smaller dimension $\mathbf{z}_n \in \mathbb{R}^M$
- We have now 2 latent variables:
- Z a N × M matrix that gives features for the users
- \mathbf{W} a $D \times M$ matrix that gives features for the movies

$$x_{dn} \approx \mathbf{w}_d^T \mathbf{z}_n$$

 We can add a regularizer and minimize the following cost:

$$\mathcal{L}(\mathbf{W}, \mathbf{Z}) = \frac{1}{2} \sum_{n=1}^{N} \sum_{d=1}^{D} (x_{dn} - \mathbf{w}_{d}^{T} \mathbf{z}_{n})^{2}$$

$$+\frac{\lambda_w}{2}\sum_{d=1}^D \mathbf{w}_d^T \mathbf{w}_d + \frac{\lambda_z}{2}\sum_{n=1}^N \mathbf{z}_n^T \mathbf{z}_n$$

first minimizing w.r.t. Z given W and then minimizing W given Z. This is called Alternating least-squares (ALS):

$$\mathbf{z}^T \leftarrow (\mathbf{w}^T \mathbf{w} + \lambda_z \mathbf{I}_M)^{-1} \mathbf{w}^T \mathbf{x}$$
$$\mathbf{w}^T \leftarrow (\mathbf{z}^T \mathbf{z} + \lambda_w \mathbf{I}_M)^{-1} \mathbf{z}^T \mathbf{x}^T$$

- Complexity: $O(DNM^2 + NM^3) \rightarrow O(DNM^2)$

tion of Gaussian mixture model
$$Y = \prod_{n=1}^{N} [p(\mathbf{x}_n | \mathbf{r}_n, \boldsymbol{\mu}, \boldsymbol{\Sigma}) p(\mathbf{r}_n | \boldsymbol{\pi})] = \prod_{n=1}^{N} [p(\mathbf{x}_n | \mathbf{r}_n, \boldsymbol{\mu}, \boldsymbol{\Sigma}) p(\mathbf{r}_n | \boldsymbol{\pi})] = \prod_{n=1}^{K} [(\mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k))^T nk] \prod_{k=1}^{K} \prod_{d \in On} \mathcal{N}(\mathbf{x}_{dn} | \mathbf{w}_d^T \mathbf{z}_n, I) \times \prod_{n=1}^{N} \mathcal{N}(\mathbf{z}_n | 0, \frac{1}{\lambda_z} I) = \prod_{l \in On} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod_{l \in I} \mathcal{N}(\mathbf{w}_d | 0, \frac{1}{\lambda_w} I) = \prod$$

Since many ratings are missing we cannot normalize the data. A solution is to add offset

$$\frac{1}{2} \sum_{n=1}^{N} \sum_{d \in O_n} (x_{dn} - \mathbf{w}_d^T \mathbf{z}_n - w_{0d} - z_{0n} - \mu)^2$$

19 Singular Value Decomposition

- Matrix factorization method

$$\mathbf{X} = \mathbf{U}\mathbf{S}\mathbf{V}^T$$

- \mathbf{U} is an $D \times D$ matrix
- **V** is an $N \times N$ matrix - S is a non-negative diagonal matrix of size $D \times N$ which are called singular values appearing in a descending order.
- Columns of U and V are the left and right
- singular vectors respectively. Assuming D < N we have

Assuming
$$D < N$$
 we have
$$\mathbf{X} = \sum_{d}^{D} s_{d} \mathbf{u}_{d} \mathbf{v}_{d}^{T}$$

This tells you about the spectrum of \boldsymbol{X} where higher singular vectors contain the low-frequency information and lower singular values contain the high-frequency information.

Let's now truncate these matrices

$$\mathbf{X} \approx \mathbf{U}_{tr} \mathbf{S}_{tr} \mathbf{V}_{tr}^*, \mathbf{T}_{tr} \approx \mathbf{U}_{tr} \mathbf{S}_{tr}, \mathbf{T}_{te} \approx \mathbf{X}_{te} \mathbf{V}_{tr}$$

with \mathbf{T}_{tr} the reduced feature set of \mathbf{X} and \mathbf{T}_{te} the reduced feature set of \mathbf{X}_{te}

20 Principal Componement Analysis

- PCA is a dimensionality reduction method and a method to decorrelate the data $\mathbf{X} \approx \tilde{\mathbf{X}} = \mathbf{W}\mathbf{Z}^T$ such that columns of \mathbf{W} are
- orthogonal. If the data is zero mean

$$\Sigma = \frac{1}{N} \mathbf{X} \mathbf{X}^T \Rightarrow \mathbf{X} \mathbf{X}^T = \mathbf{U} \mathbf{S}^2 \mathbf{U}^T$$
$$\Rightarrow \mathbf{U}^T \mathbf{X} \mathbf{X}^T \mathbf{U} = \mathbf{U}^T \mathbf{U} \mathbf{S}^2 \mathbf{U}^T \mathbf{U} = \mathbf{S}^2$$

- Thus the columns of matrix U are called the principal components and they decorrelate
- the covariance matrix. Using SVD, we can compute the matrices in the following way

$$W = US^{1/2}, Z = VS^{1/2}$$

21 Belief Propagation

- the goal is to learn inference of the latent variables
- using belief propagation Given a directed acyclic graph G and parameters θ , a Bayesian network defines the joint distribution as follows

$$p_{\theta}(\mathbf{x}) = \prod_{k=1}^{K} p_{\theta}(x_k | \text{parents}_k)$$

- Reduce complexity using bayesian rule
- - Assign each φ_l, to a cluster C_i
- 2. Initial potential $\psi_i(\mathbf{C}_i) = \prod_{\phi_k \in \mathbf{C}_i} \phi_k$ 3. Initial all messages to 1

4. Repeat: select and edge
$$(i,j)$$
 and:
$$m_{i \to j}(\mathbf{S}_{i,j}) = \sum_{\mathbf{C}_i - \mathbf{S}_{i,j}} \psi_i \times \prod_{k \in (\mathcal{N}_i - j)} m_{k \to i}$$

- Belief: $w_i(\mathbf{C}_i) = \psi_i \times \prod_{k \in \mathcal{N}_i} \delta_{k \to i}$

22 Multi-Layer Perceptron (MLP)

- There are nonlinear function classes, whose conveninent layered structure leads to efficient computation of gradients.
- Known as feed-forward neural network $-\mathbf{z}_{n}^{(k)}$ is the k'th hidden vector $\mathbf{a}_{n}^{(k)}$ is the corresponding activation

- There are a total of K layers

$$a_{mn}^{(k)} = (\mathbf{w}_{m}^{(k)})^{T} \mathbf{z}_{n}^{(k-1)}, \quad z_{mn}^{(k)} = h(a_{mn}^{(k)})$$

- A 1-layer MLP is simply a generalization of
- linear/logistic regression - $\mathbf{B}^{(k)}$ a matrix with rows $(\mathbf{w}_m^{(k)})^T$

$$\begin{aligned} \mathbf{a}_n^{(k)} &= \mathbf{B}^{(k)} \mathbf{z}_n^{(k-1)}, \quad \mathbf{z}_n^{(k)} = h(\mathbf{a}_n^{(k)}) \\ &- \text{ thus we have the input-output relationship} \end{aligned}$$

$$\hat{y}_n = g((\mathbf{w}^{(K-1)})^T * h(\mathbf{B}^{(K-2)} * h(* \dots * h(\mathbf{B}^{(1)} * \mathbf{with} g)))$$
 with $g()$ the link function

- We learn parameters B using stochastic
- gradient-descent Frequently used transfer function

$$g(a) = \tanh(a) = \frac{e^a - e^{-a}}{e^a + e^{-a}}$$

- Backpropagation: technic to compute the gradient in time linear in the number of training points and the number of weights and is nothing else than a chain rule of differential calculus.
- The key-idea is to express the derivatives in terms of activations and hidden variables.
- 1. Forward: compute $\mathbf{a}_n^{(k)}$ and $\mathbf{z}_n^{(k)}$ 2. Backward: compute $\delta_n^{(k)} = d\mathcal{L}/d\mathbf{a}_n^{(k)}$ using

$$\boldsymbol{\delta}_n^{(k-1)} = diag[(h'(\mathbf{a}_n^{(k-1)})](\mathbf{B}^{(k)})^T \boldsymbol{\delta}_n^{(k)}$$

 $\delta_n^K = \mathbf{a}_n^K - t$: 3. Compute $d\mathcal{L}/d\mathbf{B}^{(k)}$ using

$$\frac{d\mathcal{L}}{d\mathbf{B}^{(k)}} = \sum_{n} \delta_{n}^{(k)} (\mathbf{z}_{n}^{(k)})^{T}$$

23 Gaussian Process (GP)

- GP are a family of statistical distributions in which time plays a role and for which any finite linear combination of samples has a joint
- Gaussian distribution non-parametric method (rather uses latent variables) that compute a probability dist, over
- predictions They can be completely defined by their second-order statistics which means that if they have mean zero, then defining the covariance function completely defines the process
- behaviour. Let us place a probabilistic prior shape on the
- approximation of a function. A GP process defines a prior over function f

$$p(\mathbf{f}|X) = \mathcal{N}(\mathbf{f}|\mathbf{0}, K(X))$$

K(X) defines shape and prior knowledge about

$$p(\mathbf{f}|\mathbf{y}, \mathbf{X}_*, \mathbf{X}) \sim \mathcal{N}(\mu', \sigma')$$

 $\mu' = K(\mathbf{X}_*, \mathbf{X})[K(\mathbf{X}, \mathbf{X}) + \sigma_v^2 I]^{-1}\mathbf{y}$

 $\sigma' = K(\mathbf{X}_*, \mathbf{X}_*) - K(\mathbf{X}_*, \mathbf{X})[K(\mathbf{X}, \mathbf{X}) + \sigma_n^2 I]K(\mathbf{X},$

with
$$\sigma_n$$
 the variance of the noise Quadratic kernel

$k(\mathbf{x}_n, \mathbf{x}_m) = (1 + \mathbf{x}_n^T \mathbf{x}_m)^2$

- 24 Decision Trees (DT) - A decision tree is a structure in which each internal node represents a test on an attribute and each branch represents the outcome of the test and each leaf represents a class label.
- Fast to train and fast to make predictions Efficient for very high dimensional feature
- Lack of smoothness and high variance (overfitting) Goal: find a split (k, τ) that minimizes an

spaces and very large amounts of training data

impurity measure at the leaves - Find best feature to split on - Find best threshold

25 Random Forests (RF)

- RF correct the overfitting bad "habit" of DTs Training: Learn M trees on different subsets
- (random) of training data

Prediction: Average of prediction of each tree
 Variance of the model averaging:

$$z_1 = f_1 \quad \Rightarrow \quad z_M = \frac{1}{M} \, \sum_{i=1}^M f_i$$

$$\begin{split} z_1 &= f_1 \quad \Rightarrow \quad z_M = \frac{1}{M} \sum_{i=1}^M f_i \\ V(z_1) &= \sigma^2 \quad \Rightarrow \quad V(z_M) = \frac{1}{M} \sigma^2 + \rho \frac{M-1}{M} \sigma^2 \end{split}$$

- Variance reduction ratio:

$$\frac{V(z_1)}{V(z_M)} = \frac{M}{1+\rho(M-1)}$$

- 2 techniques for decorrelating trees:
 - Bagging: Randomize training data
 Randomized feature selection