Optimization

- Many more complex models in statistics and machine learning do not have analytic solutions to ML estimators.
- In most models parameters are learned by some numerical optimization technique.

$\min_{\theta} F(\theta)$

- How many minima are there?
- How do we find optimal θ ?
- Are we guaranteed to find the global optimum θ^* , rather just a local one?
- How efficiently can we solve for θ ?
- What if there are constraints?

Constrained Optimization

Optimization problems with constraints, e.g.

```
 \min_{\theta \in \mathbb{R}^d} F(\theta) 
subject to g_i(\theta) \le 0 for i = 1, ..., I
h_j(\theta) = 0 for j = 1, ..., J
```

where g_i enforce inequality constraints and h_j equality constraints.

Can write this succinctly:

 $\min_{\theta \in \mathbb{R}^d} F(\theta)$ subject to $g(\theta) \leq 0$ $h(\theta) = 0$

where $g : \mathbb{R}^d \to \mathbb{R}^I$ is a vector-valued function with $g(\theta)_i = g_i(\theta)$. Similarly $h(\theta) : \mathbb{R}^d \to \mathbb{R}^J$. $x \leq y$ iff $x_i \leq y_i \forall i$.

These problems are called programmes.

Constrained Optimization

 $\min_{\theta \in \mathbb{R}^d} F(\theta)$ subject to $g(\theta) \leq 0$ $h(\theta) = 0$

- We can enforce constraints by using Lagrange multipliers or dual variables λ ∈ ℝ^I and κ ∈ ℝ^J.
- The optimization problem can be written as a mini-max optimization of the Lagrangian:

$$\min_{\theta} \max_{\lambda \succeq 0, \kappa} \mathcal{L}(\theta, \lambda, \kappa) = \min_{\theta} \max_{\lambda \succeq 0, \kappa} F(\theta) + \lambda^{\top} g(\theta) + \kappa^{\top} h(\theta)$$

• Intuition: For any θ , we have:

 $\max_{\lambda \succeq 0, \kappa} \mathcal{L}(\theta, \lambda, \kappa) = \begin{cases} +\infty & \text{if there is some unsatisfied constraint,} \\ F(\theta) & \text{if all constraints are satisfied.} \end{cases}$

So the outer minimization over θ results in the same optimization problem.

Convex Optimization

• A function $f : \mathbb{R}^d \to \mathbb{R}$ is **convex** if

 $f(\alpha x + (1 - \alpha)y) \le \alpha f(x) + (1 - \alpha)f(y)$

for all $x, y \in \mathbb{R}^d$, $\alpha \in [0, 1]$.

- For smooth functions: Equivalent to 2nd derivative (Hessian) being positive semidefinite.
- A programme is a convex programme if:
 - $F(\theta)$ is convex,
 - $g_i(\theta)$ is convex for each *i*,
 - $h(\theta) = A\theta + b$ is affine.
- Examples: linear, quadratic, semidefinite programming.
- Convex programmes have a unique minimum (typically), which can be efficiently found.



Convex Duality

- Say the minimum is p^* , and occurred at θ^* .
- The dual programme inverts the order of max and min:

 $p^* = \min_{\theta} \max_{\lambda \succeq 0, \kappa} \mathcal{L}(\theta, \lambda, \kappa) \geq \max_{\lambda \succeq 0, \kappa} \min_{\theta} \mathcal{L}(\theta, \lambda, \kappa) = d^*$

where the dual optimum is d^* .

- Karush-Kuhn-Tucker Theorem: Subject to regularity conditions, a solution θ* is the optimal solution of a convex programme, if and only if there are λ* and κ* (the dual optimal solution) such that:
 - ▶ Primal feasible: $g(\theta^*) \leq 0$, $h(\theta^*) = 0$.
 - Dual feasible: $\lambda^* \succeq 0$.
 - $(\theta^*, \lambda^*, \kappa^*)$ is a saddle point of \mathcal{L} : For every $\theta, \lambda \succeq 0, \kappa$, we have

$$\mathcal{L}(\theta^*, \lambda, \kappa) \leq \mathcal{L}(\theta^*, \lambda^*, \kappa^*) \leq \mathcal{L}(\theta, \lambda^*, \kappa^*)$$

- Complementary slackness: For every *i*,

 $\lambda_i^* g_i(\theta^*) = 0$

Linear Classification

- ► A dataset with {+1, -1} labels is **linearly separable** if there is a hyperplane separating two classes.
- Typically there will be an infinite number of such separating hyperplanes.



Maximum Margin Classification

- ► Good choice of separating hyperplane: one with large margin.
- Such a hyperplane will be defined by a number of data vectors close to the boundary—the support vectors, leading to a method called support vector machines.



Support Vector Machines

A hyperplane can be parametrized as:

 $g(x) = a + b^{\top}x = 0$

with the classification given by sign(g(x)).

- Distance and classification of a point x_i from hyperplane is $g(x_i)/||b||$.
- Multiplying a and b by c > 0 does not affect result. Rescale such that margin (closest distance of data vectors to hyperplane) is 1/||b||.



 $y_i(a + b^{\top} x_i) / ||b|| \ge 1 / ||b||$ $y_i(a + b^{\top} x_i) \ge 1$

Constrained optimization problem to solve for *a*, *b*:

Support Vector Machines

• Introduce Lagrange multipliers $\lambda_i \ge 0$ to enforce constraints:

$$\min_{a,b} \max_{\lambda \succeq 0} \mathcal{L}(a,b,\lambda) = \frac{1}{2} \|b\|^2 + \sum_{i=1}^n \lambda_i (1 - y_i(a + b^\top x_i))$$

KKT optimality conditions:

Zero derivatives:
$$\nabla_a \mathcal{L}(a^*, b^*, \lambda^*) = -\sum_{i=1}^n \lambda_i^* y_i = 0$$
 $\nabla_b \mathcal{L}(a^*, b^*, \lambda^*) = b^* - \sum_{i=1}^n \lambda_i^* y_i x_i = 0$ Primal feasibility: $y_i(a^* + (b^*)^\top x_i) \ge 1$ Dual feasibility: $\lambda_i^* \ge 0$ Complementary slackness: $\lambda_i^* (1 - y_i(a^* + (b^*)^\top x_i)) = 0$

Support Vector Machines

Substituting optimal a* and b* into Lagrangian leads to the dual optimization problem:

$$\max_{\lambda} \sum_{i=1}^{n} \lambda_{i} - \frac{1}{2} \sum_{i,j=1}^{n} \lambda_{i} \lambda_{j} y_{i} y_{j} (x_{i})^{\top} (x_{j})$$

subject to
$$\sum_{i=1}^{n} \lambda_{i} y_{i} = 0$$
$$\lambda \succeq 0$$

A quadratic programme. Standard solvers can be used to find optimal λ^* in $O(n^3)$ cost.

- Those vectors with $\lambda_i > 0$ are called **support vectors**.
- Complementary slackness implies that if x_i does not lie on boundary, then $\lambda_i = 0$, i.e. not a support vector.
- Discriminant function is

$$g(x) = a^* + \sum_{i=1}^n \lambda_i^* y_i x_i^\top x$$

where a^* can be solved by noting that $y_j g(x_j) = 1$ for a support vector x_j .

For non-linearly separable datasets, we can allow for margin violations

$$\xi_i = \begin{cases} 1 - y_i(a + b^{\top} x_i) & \text{if margin violated,} \\ 0 & \text{if not violated.} \end{cases}$$
$$= \max(0, 1 - y_i(a + b^{\top} x_i))$$

Penalizing violations by their magnitude,

$$\min_{\substack{a,b,\xi \\ a,b,\xi}} \frac{1}{2} \|b\|^2 + C \sum_{i=1}^n \xi_i$$

subject to $y_i(a + b^\top x_i) \ge 1 - \xi_i$
 $\xi_i \ge 0$

where *C* is a tuning parameter.



• Introduce Lagrange multipliers $\lambda_i \ge 0$, $\gamma_i \ge 0$ to enforce constraints:

$$\mathcal{L}(a,b,\xi,\lambda,\gamma) = \frac{1}{2} \|b\|^2 + C \sum_{i=1}^n \xi_i + \sum_{i=1}^n \lambda_i (1 - \xi_i - y_i(a + b^\top x_i)) - \sum_{i=1}^n \gamma_i \xi_i$$

KKT optimality conditions:

 $\nabla_a \mathcal{L}(a^*, b^*, \xi^*, \lambda^*, \gamma^*) = -\sum_i \lambda_i^* y_i = 0$ Zero derivatives: $\nabla_b \mathcal{L}(a^*, b^*, \xi^*, \lambda^*, \gamma^*) = b^* - \sum_{i=1}^{n} \lambda_i^* y_i x_i = 0$ $\nabla_{\mathcal{E}_i} \mathcal{L}(a^*, b^*, \xi^*, \lambda^*, \gamma^*) = C - \lambda_i^* - \gamma_i^* = 0$ $y_i(a^* + (b^*)^\top x_i) > 1 - \xi_i^*$ **Primal feasibility:** $\xi_i^* \geq 0$ **Dual feasibility:** $\lambda_i^* \geq 0$ $\gamma_i^* \geq 0$ Complementary slackness: $\lambda_i^*(1 - \xi_i^* - y_i(a^* + (b^*)^\top x_i)) = 0$ $\gamma_i^*\xi_i^*=0$

207

Setting derivatives of primal variables to zero leads to the dual programme:

$$\max_{\lambda} \sum_{i=1}^{n} \lambda_{i} - \frac{1}{2} \sum_{i,j=1}^{n} \lambda_{i} \lambda_{j} y_{i} y_{j} (x_{i})^{\top} (x_{j})$$

subject to
$$\sum_{i=1}^{n} \lambda_{i} y_{i} = 0$$
$$0 \leq \lambda \leq C$$

Only difference is the **box constraint** on $\lambda_i \in [0, C]$.

From primal programme, we can first minimize over ξ_i 's, leading to an unconstrained convex programme:

$$\min_{a,b} \quad \frac{1}{2} \|b\|^2 + C \sum_{i=1}^n \max(0, 1 - y_i(a + b^\top x_i))$$

 Interpretation: regularized empirical risk minimization with the hinge loss.



Support Vector Machines – Discussion

- Multiclass classification: If there are more than two classes, there are multiclass generalizations of the SVM.
- A simple practical idea: treat a multiclass problem as multiple binary classification problems.
 - One-vs-one: train K(K 1) binary SVMs, for each pair of classes. At test time, predict class that got the most votes.
 - One-vs-rest: train *K* binary SVMs, one for each class vs all other classes. At test time, predict class with largest discriminant value $a_k + b_k^{\top} x$.
- Optimization for large scale problems:
 - Standard quadratic programme solvers not scalable.
 - Sequential minimal optimization (SMO): iterative solve pairs of λ_i 's.
 - Pegasos : stochastic gradient descent on regularized hinge loss objective.
- \blacktriangleright L₂ regularization controls overfitting.
- Not probabilistic and cannot produce uncertainty estimates.
- Statistical learning theory foundations.
- Further readings:
 - Bishop, Chapter 6.
 - Christopher Burgess, A Tutorial on Support Vector Machines for Pattern Recognition. 1998.

Nonlinear Methods

- Decision boundaries and regression functions often need to be nonlinear.
- One general approach: transform data $x \mapsto \phi(x)$.
- A global approach. Decisions and optimal parameters depend on whole training dataset.
- Alternative approach:
 p(Y = 1 | X = x) or f(x) depends
 only on data cases in local
 neighbourhood of x.



Local Methods



k-Nearest Neighbours

- A simple, local, nonlinear, non-model-based, method.
- Prediction at a data vector x is simply determined by the k nearest neighbours ne_k(x) of x among the training set.
- Classification: predict the majority vote of the neighbours:

$$f^{\mathsf{kNN}}(x) = \underset{l}{\operatorname{argmax}} |\{j \in ne_k(x) : y_j = l\}|.$$

Regression: predict the average among the neighbours:

$$f^{\mathsf{kNN}}(x) = \frac{\sum_{j \in ne_k(x)} y_j}{\sum_{j \in ne_k(x)} 1}.$$



k-Nearest Neighbours

- Nearest neighbours are simple and essentially model-free methods for classification.
- Weaker modelling assumptions than e.g. LDA, Naïve Bayes and logistic regression.
- These methods are not very useful for understanding relationships between attributes and class predictions.
- As black box classification methods however, they are often perform reasonably on real life problems and provide a good benchmark.
- Can break down in high-dimensional data:
 - Effectively, partitions input space into regions each containing k data points, and prediction in each region estimated separately.
 - In a space of dimension p >>> 0, number of regions needed is R = m^p, so size of dataset needed is km^p.



Data



trainx[, 1]



trainx[, 1]



trainx[, 1]



trainx[, 1]



trainx[, 1]



trainx[, 1]



trainx[, 1]

Result of 51NN

k-Nearest Neighbour Demo – R Code I

```
library(MASS)
## load crabs data data(crabs)
ct <- as.numeric(crabs[,1])-1+2*(as.numeric(crabs[,2])-1)
## project into first two LD
cb.lda <- lda(log(crabs[,4:8]),ct)
cb.ldp <- predict(cb.lda)
x <- as.matrix(cb.ldp$x[,1:2])
y <- as.numeric(crabs[,2])-1
x <- x + rnorm(dim(x)[1]*dim(x)[2])*1.5
eqscplot(x,pch=2*y+1,col=1)</pre>
```

k <- 3

```
kNN <- function(k,x,y,gridsize=100) {</pre>
```

```
n
       <- length(v)
       <- dim(x)[2]
р
       <- sample(rep(c(TRUE,FALSE),each=n/2),n,replace=FALSE)
i
train <- (1:n)[i]
test <- (1:n) [!i]
trainx <- x[train,]</pre>
trainy <- y[train]</pre>
testx <- x[test,]</pre>
testy <- y[test]</pre>
trainn <- dim(trainx)[1]</pre>
testn <- dim(testx)[1]</pre>
gridx1 <- seq(min(x[,1]), max(x[,2]), length=gridsize)</pre>
gridx2 <- seq(min(x[,2]), max(x[,2]), length=gridsize)</pre>
gridx <- as.matrix(expand.grid(gridx1,gridx2))</pre>
gridn <- dim(gridx)[1]</pre>
# calculate distances, smart and intelligently.
trainxx <- t((trainx*trainx) %*% matrix(1,p,1))</pre>
```

k-Nearest Neighbour Demo – R Code II

```
testxx <- (testx*testx) %*% matrix(1,p,1)</pre>
gridxx <- (gridx*gridx) %*% matrix(1,p,1)</pre>
testtraindist <- matrix(1,testn,1) %*% trainxx +</pre>
                  testxx %*% matrix(1,1,trainn) -
                  2*(testx %*% t(trainx))
gridtraindist <- matrix(1,gridn,1) %*% trainxx +</pre>
                  gridxx %*% matrix(1,1,trainn) -
                  2*(gridx %*% t(trainx))
# predict
testp <- numeric(testn)</pre>
gridp <- numeric(gridn)</pre>
for (j in 1:testn) {
  nearestneighbors <- order(testtraindist[j,])[1:k]</pre>
  testp[j] <- mean(trainy[nearestneighbors])</pre>
for (j in 1:gridn) {
  nearestneighbors <- order(gridtraindist[j,])[1:k]</pre>
  gridp[j] <- mean(trainy[nearestneighbors])</pre>
predy <- as.numeric(testp>.5)
plot(trainx[,1],trainx[,2],pch=trainy*3+1,col=4,lwd=.5)
points(testx[,1],testx[,2],pch=testy*3+1,col=2+(predy==testy),lwd=3)
contour(gridx1, gridx2, matrix(gridp, gridsize, gridsize),
      levels=seq(.1,.9,.1),lwd=.5,add=TRUE)
contour(gridx1, gridx2, matrix(gridp, gridsize, gridsize),
      levels=c(.5),lwd=2,add=TRUE)
```

}

Asymptotic Performance of 1NN

- ▶ Let $(x_i, y_i)_{i=1}^n$ be training data where $x_i \in \mathbb{R}^p$ and $y_i \in \{1, 2, ..., K\}$.
- We define

$$\widehat{y}_{\mathsf{Bayes}}\left(x\right) = \underset{l \in \{1, \dots, K\}}{\arg \max} \pi_{l} f_{l}\left(x\right)$$

and

$$\widehat{y}_{1NN}(x) = y$$
 (nearest neigbour of x).

► The (optimal) Bayes risk and 1NN risk are:

$$\begin{array}{lll} R_{\mathsf{Bayes}} & = & \mathbb{E}\left[\mathbb{I}\left(Y \neq \widehat{y}_{\mathsf{Bayes}}\left(X\right)\right)\right] \\ R_{\mathsf{1NN}} & = & \mathbb{E}\left[\mathbb{I}\left(Y \neq \widehat{Y}_{\mathsf{1NN}}\left(X\right)\right)\right] \end{array}$$

• As $n \to \infty$, we have the following powerful result

$$R_{\mathsf{Bayes}} \leq R_{\mathsf{1NN}} \leq 2R_{\mathsf{Bayes}} - rac{K}{K-1}R_{\mathsf{Bayes}}^2.$$

K-Nearest Neighbours – Discussion

- kNN is sensitive to distances: normalize data and find suitable metric.
- Choice of k important: controls flexibility of model.
- Computational cost of kNN is very high.
 - Need to store **all** training data.
 - Need to compare each test data vector to all training data.
 - Need a lot of data in high dimensions.
- Mitigation techniques:
 - Compute approximate nearest neighbours, using kd-trees, cover trees, random forests.
 - Apply K-means to data in each class, to reduce size of data (need to use large K).

Non-linear Problems

- Linear methods (PCA, LDA, linear and logistic regression) are simple and effective techniques to learn from data "to first order".
- To capture more intricate information from data, flexible, non-linear methods are often needed.
 - Explicit non-linear transformations $x \mapsto \phi(x)$.
 - Local methods like kNN.
- Kernel methods: introduce non-linearities through implicit non-linear transforms, often local in nature.



Back to the soft-margin SVM. The dual objective is:

$$\max_{\lambda} \quad \sum_{i=1}^{n} \lambda_{i} - \frac{1}{2} \sum_{i,j=1}^{n} \lambda_{i} \lambda_{j} y_{i} y_{j} \phi(x_{i})^{\top} \phi(x_{j}) \quad \text{subject to} \quad \begin{cases} \sum_{i=1}^{n} \lambda_{i} y_{i} = 0\\ 0 \leq \lambda \leq C \end{cases}$$

Suppose p = 2, and we would like to introduce quadratic non-linearities,

$$\phi(x_i) = (1, \sqrt{2}x_{i1}, \sqrt{2}x_{i2}, x_{i1}^2, x_{i2}^2, x_{i1}x_{i2})^{\mathsf{T}}$$

Then

$$\phi(x_i)^{\top} \phi(x_j) = 1 + 2x_{i1}x_{j1} + 2x_{i2}x_{j2} + x_{i1}^2x_{j1}^2 + x_{i2}^2x_{j2}^2 + x_{i1}x_{i2}x_{j1}x_{j2}$$
$$= (1 + x_i^{\top}x_j)^2$$

- Since only dot-products are needed in the objective function, non-linear transform need not be computed explicitly!
- Generally, *m*-order interactions can be implemented simply by $\phi(x_i)^{\top}\phi(x_j) = (1 + x_i^{\top}x_j)^m$. This is called a **polynomial kernel**.

► The **Gram matrix** is the matrix of dot-products, $B_{ij} = \phi(x_i)^\top \phi(x_j)$.

$$B = \begin{pmatrix} -\phi(x_1)^\top & -\\ \vdots \\ -\phi(x_i)^\top & -\\ \vdots \\ -\phi(x_n)^\top & - \end{pmatrix} \times \begin{pmatrix} | & | & | \\ \phi(x_1) & \cdots & \phi(x_j) & \cdots & \phi(x_n) \\ | & | & | \end{pmatrix}$$

- Since $B = \Phi \Phi^{\top}$, it is symmetric and positive semidefinite.
- ► The Gram matrix is sufficient for training the soft-margin SVM.

$$\max_{\lambda} \sum_{i=1}^{n} \lambda_{i} - \frac{1}{2} \sum_{i,j=1}^{n} \lambda_{i} \lambda_{j} y_{i} y_{j} B_{ij} \qquad \text{subject to} \qquad \begin{cases} \sum_{i=1}^{n} \lambda_{i} y_{i} = 0\\ 0 \leq \lambda \leq C \end{cases}$$

- A **kernel** is a function $\kappa : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ such that:
 - Symmetric: $\kappa(x, x') = \kappa(x', x)$.
 - ► Positive semidefinite: given any finite set $\{x_i\}_{i=1}^n \subset \mathcal{X}$, the matrix $B \in \mathbb{R}^{n \times n}$ with entries $B_{ij} = \kappa(x_i, x_j)$ is positive definite. Equivalently, for any $c \in \mathbb{R}^n$,



Mercer's Theorem: if κ is continuous, symmetric and positive semidefinite, then there is a function φ : X → H into a Hilbert space H with inner product ⟨·, ·, ⟩ such that

$$\kappa(x, x') = \langle \phi(x), \phi(x') \rangle$$

 $\kappa(x, x') = \langle \phi(x), \phi(x') \rangle$

We do not need to compute the features ever—the Gram matrix is sufficient for learning and prediction. The discriminant function (absorbing *a* into *b*) is

$$g(x) = \sum_{i=1}^{n} \lambda_i^* y_i \phi(x_i)^\top \phi(x) = \sum_{i=1}^{n} \lambda_i^* y_i \kappa(x_i, x)$$

- ► The function \u03c6 can be interpreted as non-linear features of our data vectors x ∈ \u03c8.
- Generally, the Hilbert space can be infinite-dimensional, so we are effectively computing an infinite number of features of our data, and learning a SVM based on all features.
- There are an infinite number of parameters in the SVM—a nonparametric method.
- The L_2 regularization of SVMs is very important to prevent overfitting.

Examples of Kernels

Polynomial kernel:

$$\kappa(x, x') = (1 + x^\top x')^m$$

Gaussian, radial-basis function (RBF), or squared-exponential kernel:

$$\kappa(x, x') = \exp\left(-\frac{1}{2}\|x - x'\|_M^2\right)$$

This leads to a discriminant function of form

$$g(x) = \sum_{i=1}^{n} \lambda_i^* y_i \exp\left(-\frac{1}{2} ||x_i - x||_M^2\right)$$

A local method very similar to kNN.

• If κ_1 and κ_2 are both kernels, then so are kernels defined by

$$\kappa_3(x, x') = \kappa_1(x, x') + \kappa_2(x, x')$$

$$\kappa_4(x, x') = \kappa_1(x, x') \times \kappa_2(x, x')$$

Kernel SVM Demo

```
library(MASS)
library(e1071)
## load crabs data, project onto LD space, add noise.
data(crabs)
ct <- as.numeric(crabs[,1])-1+2*(as.numeric(crabs[,2])-1)</pre>
cb.lda <- lda(log(crabs[,4:8]),ct)</pre>
cb.ldp <- predict(cb.lda)</pre>
x <- as.matrix(cb.ldp$x[,1:2])</pre>
y <- as.numeric(crabs[,2])-1</pre>
x < -x + rnorm(dim(x)[1] * dim(x)[2]) * 1.5
gridsize <- 100
xlim <- c(min(x[,1]), max(x[,1]))
ylim <- c(min(x[,2]), max(x[,2]))
gridx1 <- seg(xlim[1], xlim[2], length=gridsize)</pre>
gridx2 <- seg(ylim[1], ylim[2], length=gridsize)</pre>
gridx <- as.matrix(expand.grid(gridx1,gridx2))</pre>
gridn <- dim(gridx)[1]</pre>
plot(x,pch=2*y+1,col=1,xlim=xlim,ylim=ylim)
       <- length(y)
n
р
       <- dim(x)[2]
i
       <- sample(rep(c(TRUE,FALSE),each=n/2),n,replace=FALSE)
train <- (1:n)[i]
test <- (1:n) [!i]
trainx <- x[train,]</pre>
trainy <- y[train]</pre>
testx <- x[test,]</pre>
testy <- y[test]</pre>
svmdemo <- function(kernel,gamma=1,coef0=0,cost=1,degree=3) {</pre>
  model <- svm(trainx,trainy,kernel=kernel,gamma=gamma,coef0=coef0,degree=degree,cost=cost)</pre>
  gridp <- predict(model,gridx)</pre>
  predy <- as.numeric(predict(model,testx)>.5)
  plot(trainx[,1],trainx[,2],pch=trainy*3+1,col=4,lwd=.5,xlim=xlim,ylim=ylim)
  points(testx[,1],testx[,2],pch=testy*3+1,col=2+(predy==testy),lwd=3)
  contour(gridx1,gridx2,matrix(gridp,gridsize,gridsize),levels=seg(.1,.9,.1),lwd=.5,add=TRUE)
  contour(gridx1,gridx2,matrix(gridp,gridsize,gridsize),levels=c(.5),lwd=2,add=TRUE)
}
```

233

Kernel Methods – Discussion

- The kernel method allows for very flexible and powerful machine learning models.
- Kernels can be defined over much more complex structures than vectors, e.g. graphs, strings.
- Can be hard to interpret.
- $O(n^3)$ computation and $O(n^2)$ memory cost can be prohibitive.
- Further readings:
 - Bishop, Chapter 6.
 - Christopher Burgess, A Tutorial on Support Vector Machines for Pattern Recognition. 1998.
 - Rasmussen and Williams, Gaussian Processes for Machine Learning. 2006.