#### **STAT540**

Lecture 18: March 16<sup>th</sup> 2015

#### Supervised learning II

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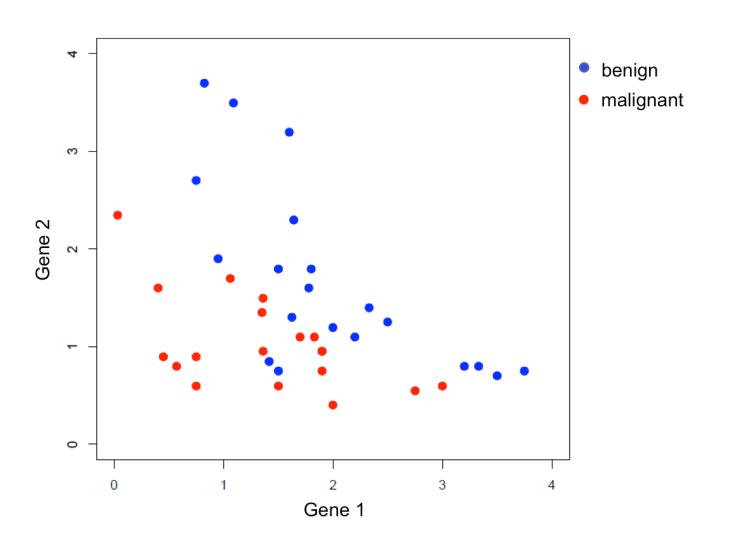
Center for Molecular Medicine and Therapeutics

#### Outline

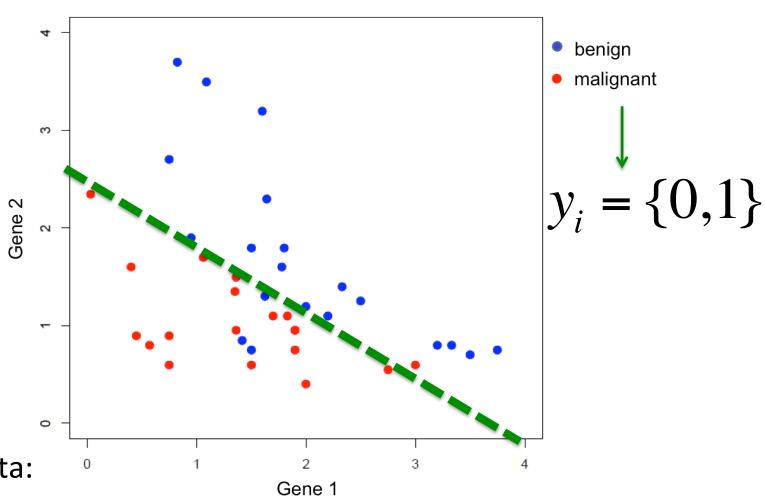
- Binary classification:
  - Logistic regression (linear)
  - Support vector machines ("hyper-linear")

- Complexity and model selection:
  - Overfitting
  - Cross-validation

#### Binary classification task



#### The decision boundary



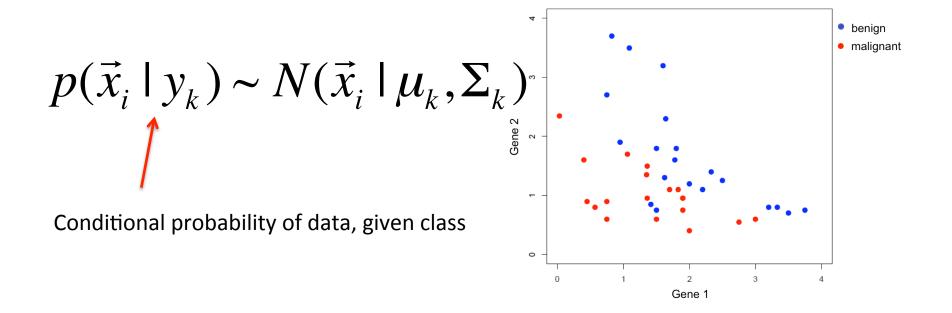
Training data:

$$D_T = \{(\vec{x}_1, y_1), ..., (\vec{x}_n, y_n)\}$$

#### Linear Disciminant Analysis

(by Fisher in 1936)

 Assume that gene 1 and gene 2 are normally distributed in each class



#### Linear Disciminant Analysis

Prior class probability  $\frac{p(y_i = 0 \mid \vec{x}_i)}{p(y_i = 1 \mid \vec{x}_i)} > 1 \Leftrightarrow \log(\frac{N(\vec{x}_i \mid \mu_1, \Sigma)p(y_i = 0)}{N(\vec{x}_i \mid \mu_2, \Sigma)p(y_i = 1)}) > 0$ 

#### Linear Disciminant Analysis

Posterior probability of class

terior probability of class 
$$\frac{p(y_i = 0 \mid \vec{x}_i)}{p(y_i = 1 \mid \vec{x}_i)} > 1 \Leftrightarrow \log(\frac{N(\vec{x}_i \mid \mu_1, \Sigma)p(y_i = 0)}{N(\vec{x}_i \mid \mu_2, \Sigma)p(y_i = 1)}) > 0$$

Prior class probability

$$\Leftrightarrow \vec{a}^T \vec{x}_i + b > 0$$

where

$$\vec{a} = \Sigma^{-1}(\mu_1 - \mu_2)$$

$$\vec{b} = (\mu_1 - \mu_2)^T \Sigma^{-1}(\mu_1 - \mu_2) - \log(\frac{p(y_i = 0)}{p(y_i = 1)})$$

$$\log\left(\frac{p(y_i = 0 \mid \vec{x}_i)}{p(y_i = 1 \mid \vec{x}_i)}\right) = \log\left(\frac{p(y_i = 0 \mid \vec{x}_i)}{1 - p(y_i = 0 \mid \vec{x}_i)}\right) = \vec{a}^T \vec{x}_i + b$$

Derive the maximum likelihood estimate (MLE) for the parameters (a and b)

$$\log\left(\frac{p(c=1|\vec{x}_i)}{p(c=2|\vec{x}_i)}\right) = \log\left(\frac{p(c=1|\vec{x}_i)}{1-p(c=1|\vec{x}_i)}\right) = \vec{a}^T \vec{x}_i + b$$

#### **Data Likelihood**

$$L(D \mid \theta) = \prod_{i=1}^{n} p(y_i \mid \vec{x}_i)$$
| IID Samples

Mathematical convenience

Data Likelihood 
$$L(D \mid \theta) = \prod_{i=1}^{n} p(y_i = 0 \mid \vec{x}_i)^{1-y_i} p(y_i = 1 \mid \vec{x}_i)^{y_i}$$

To find the MLE, we need to know:  $p(y_i \mid \vec{x}_i)$  for  $y_i = \{0,1\}$ 

$$\log\left(\frac{p(y_i = 0 \mid \vec{x}_i)}{1 - p(y_i = 0 \mid \vec{x}_i)}\right) = \vec{a}^T \vec{x}_i + b$$

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$$\log\left(\frac{p(y_i = 0 \mid \vec{x}_i)}{1 - p(y_i = 0 \mid \vec{x}_i)}\right) = \vec{a}^T \vec{x}_i + b$$

$$p(y_{i} = 0 \mid \vec{x}_{i}) = \frac{e^{\vec{w}^{T} \vec{x}_{i}}}{1 + e^{\vec{w}^{T} \vec{x}_{i}}}$$

$$w=[b a]$$

$$x_{i}=[1 x_{i}]$$

$$p(y_{i} = 1 \mid \vec{x}_{i}) = \frac{1}{1 + e^{\vec{w}^{T} \vec{x}_{i}}}$$

# Logistic regression: MLE

$$L(D \mid \theta) = \prod_{i=1}^{n} p(y_i = 0 \mid \vec{x}_i)^{1-y_i} p(y_i = 1 \mid \vec{x}_i)^{y_i} = \prod_{i=1}^{n} \left(\frac{1}{1 + e^{(\vec{w}^T \vec{x}_i)}}\right)^{y_i} \left(\frac{e^{(\vec{w}^T \vec{x}_i)}}{1 + e^{(\vec{w}^T \vec{x}_i)}}\right)^{1-y_i}$$

$$\operatorname{argmax}_{w} L(D; w) = \operatorname{argmin}_{w} - L(D; w) = \operatorname{argmin}_{w} - \log L(D; w)$$

What does it mean? For all possible w's, find the one that maximizes the likelihood.

## Logistic regression: MLE

**MLE** 

$$\operatorname{argmin}_{\mathcal{W}} - \ell(D; w)$$

$$= \operatorname{argmin}_{W} - \sum_{i=1}^{n} y_{i} \log \frac{1}{1 + e^{y_{i}(\vec{w}^{T}\vec{x}_{i})}} + (1 - y_{i}) \log \frac{e^{y_{i}(\vec{w}^{T}\vec{x}_{i})}}{1 + e^{y_{i}(\vec{w}^{T}\vec{x}_{i})}}$$



Final objective function

### Optimization problem

$$f(w)$$

$$\operatorname{argmin}_{W} - \sum_{i=1}^{n} y_{i} \log \frac{1}{1 + e^{y_{i}(\vec{w}^{T}\vec{x}_{i})}} + (1 - y_{i}) \log \frac{e^{y_{i}(\vec{w}^{T}\vec{x}_{i})}}{1 + e^{y_{i}(\vec{w}^{T}\vec{x}_{i})}}$$

- We need to differentiate the objective function wrt w to get the gradient for w.
- Differentiating the objective function reveals that there is no closed-form solution for w.
- We can use an iterative procedure (e.g., gradient descent) to find the optimal w.

# Optimization problems & gradient descent

- Consider the likelihood function f(w): need to find w that minimizes the likelihood function.
- No-closed form solution for w: numerical methods that update the solution w at each
  - Gradient descent, conjugate gradient, Newtwon method

• Update form: 
$$w^{(k)} \leftarrow w^{(k-1)} - \gamma \nabla f(w)$$

# Gradient descent for logistic regression

#### Simple for loop for fining the optimal value of w

$$w^{\text{new}} = \mathbf{0}$$
 $w^{\text{old}} = \mathbf{0}$ 

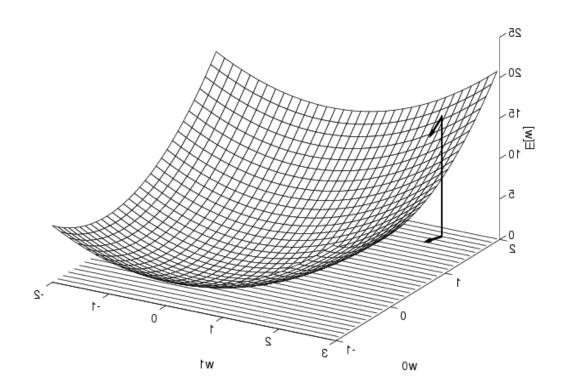
While  $||w^{\text{new}} - w^{\text{old}}||_2 > e$ 
 $w^{\text{old}} = w^{\text{new}}$ 
 $w^{\text{new}} \leftarrow w^{\text{old}} - \gamma \nabla f(w)$ 

## Summary: logistic regression

- Defined a model for quantifying data likelihood
- Simplify the likelihood if possible
- Write down an optimization problem for optimizing data likelihood given parameters
  - Use gradient descent or other more sophisticated numerical methods for estimating model parameters.

# Local vs. global minimum

Concave objective function: one optimal solution (no local minima)



Gradient of the objective function points to the direction of global minimum

# Support Vector Machines (SVMs)

- Input-output relationships may not be linear
- SVMs kernel trick: make linear model work in non-linear setting.

# What is a support vector machine?

- 1. A subset of training examples (support vectors)
- 2. A vector of weights for the support vectors ( $\vec{\alpha}$ )
- 3. A similarity function:  $K(\mathbf{x}_1, \mathbf{x}_2)$  (kernel function)

Predicting class labels for example  $x_i$ :

$$f(\vec{x}_i) = \text{sign}(\sum_j \alpha_j y_j K(\vec{x}_i, \vec{x}_j))$$

$$y_i = \{-1,1\}$$
 We are switching to -1,1 class labels for mathematical convenience

# Examples of kernels

Linear kernel: 
$$K(\vec{x}_i, \vec{x}_j) = \vec{x}_i^T \vec{x}_j$$

Polynomial kernel: 
$$K(\vec{x}_i, \vec{x}_j) = (\vec{x}_i^T \vec{x}_j)^d$$

Gaussian kernel: 
$$K(\vec{x}_i, \vec{x}_j) = \exp(-\frac{\|\vec{x}_i - \vec{x}_j\|}{\sigma})$$

# Examples of kernels

Parameters you can tune

Linear kernel: 
$$K(\vec{x}_i, \vec{x}_j) = \vec{x}_i^T \vec{x}_j$$

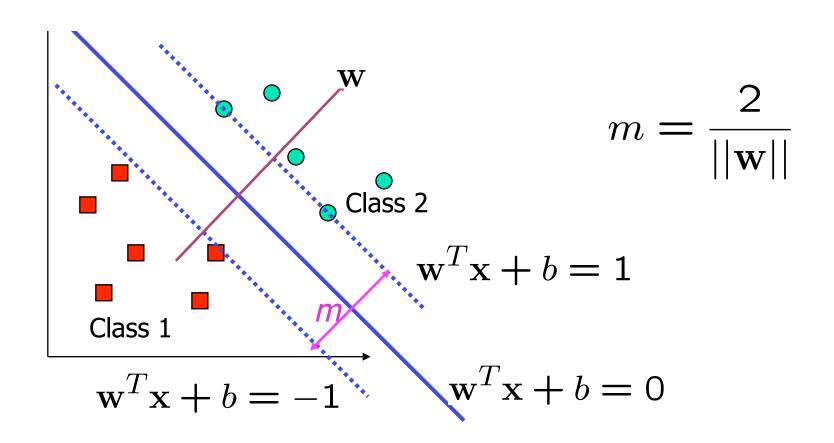
Polynomial kernel:  $K(\vec{x}_i, \vec{x}_j) = (\vec{x}_i^T \vec{x}_j)^d$ 

Gaussian kernel:  $K(\vec{x}_i, \vec{x}_j) = \exp(-\frac{||\vec{x}_i - \vec{x}_j||}{C})$ 

### Learning SVMs

- Choose a kernel function
  - How? black art or CV
- Choose the support vectors
  - How? Side effect of learning the "weights"  $\vec{lpha}$
- Choose the weights
  - Write down an objective function that maximizes
     the margin (primal form → dual form gives you the "weights")

#### Maximizing the margin



Note Distance between point and hyperplane:

$$\frac{\mid \mathbf{x}_i \cdot \mathbf{w} + b}{\parallel \mathbf{w} \parallel}$$

#### Finding the maximum margin hyperplane

- Maximize margin  $2/||\mathbf{w}||$
- Correctly classify all training data:

```
\begin{cases} \mathbf{x}_i \text{ positive } (y_i = 1) : & \mathbf{x}_i \cdot \mathbf{w} + b \ge 1 \\ \mathbf{x}_i \text{ negative } (y_i = -1) : & \mathbf{x}_i \cdot \mathbf{w} + b \le -1 \end{cases}
```

#### Finding the maximum margin hyperplane

- Maximize margin  $2/||\mathbf{w}||$
- Correctly classify all training data:

$$\begin{cases} \mathbf{x}_i \text{ positive } (y_i = 1) : & \mathbf{x}_i \cdot \mathbf{w} + b \ge 1 \\ \mathbf{x}_i \text{ negative } (y_i = -1) : & \mathbf{x}_i \cdot \mathbf{w} + b \le -1 \end{cases}$$

#### Quadratic optimization problem:

Minimize 
$$\frac{1}{2}\mathbf{w}^T\mathbf{w}$$

Subject to  $y_i(\mathbf{w} \cdot \mathbf{x}_i + b) \ge 1$  for all  $(\mathbf{x}_i, y_i)$ 

### Solving the SVM Optimization Problem

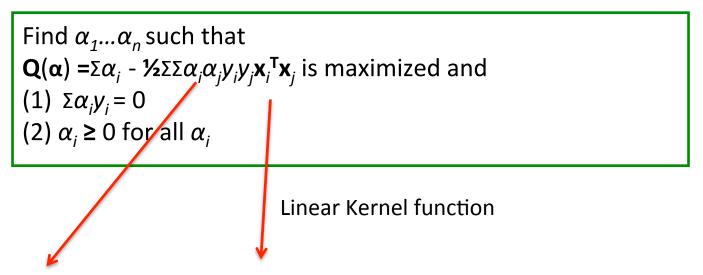
```
Find \mathbf{w} and \mathbf{b} such that \underset{\mathbf{w}}{\operatorname{argmin}}_{\mathbf{w}} \mathbf{w}^{\mathsf{T}} \mathbf{w} and for all (\mathbf{x}_{i}, y_{i}), i = 1..n: y_{i} (\mathbf{w}^{\mathsf{T}} \mathbf{x}_{i} + b) \geq 1
```

- Need to optimize a quadratic function subject to linear constraints.
- The solution involves constructing a *dual problem* where a Lagrange multiplier  $\alpha_i$  is associated with every inequality constraint in the primal (original) problem:

```
Find \alpha_1...\alpha_n such that \mathbf{Q}(\alpha) = \Sigma \alpha_i - \frac{1}{2} \Sigma \Sigma \alpha_i \alpha_j y_i y_j \mathbf{x}_i^\mathsf{T} \mathbf{x}_j is maximized and (1) \Sigma \alpha_i y_i = 0 (2) \alpha_i \ge 0 for all \alpha_i
```

### Solving the SVM Optimization Problem

Dual problem:



"support vector" weights

$$K(\vec{x}_i, \vec{x}_j) = \vec{x}_i^T \vec{x}_j$$

# **SVMs** summary

#### Pros

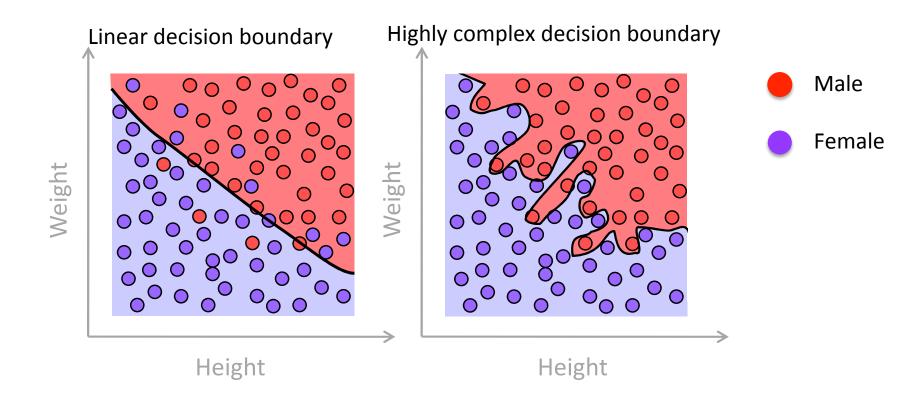
- Lots of publically available software.
- Kernel based framework is very powerful, flexible.
- Works well in practice, but need to have good number of training examples.

#### Cons

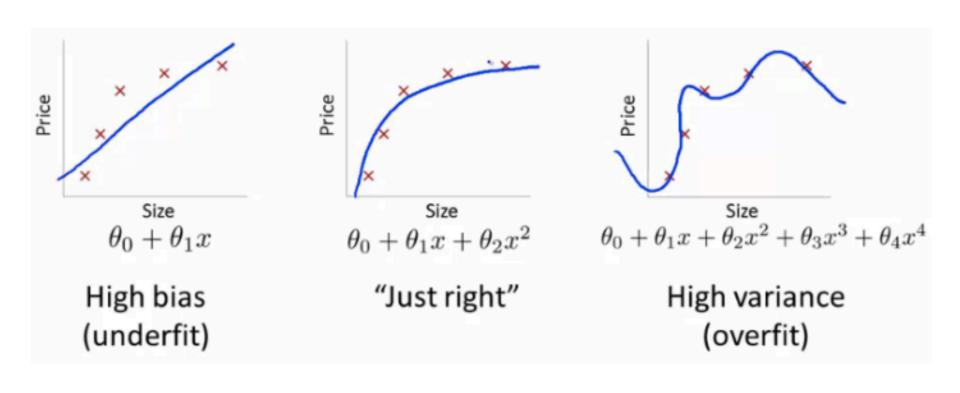
- Need to define suitable kernel (not hard to do, just use linear kernel).
- Hard optimization problem: training could take a long time depending on size of your problem.

#### **Overfitting**

If we allow very complicated predictors, we could overfit the training data:



# Overfitting example: polynomial of degree k for $k=\{1,..,4\}$ (regression problem)



#### Ockam's Razor

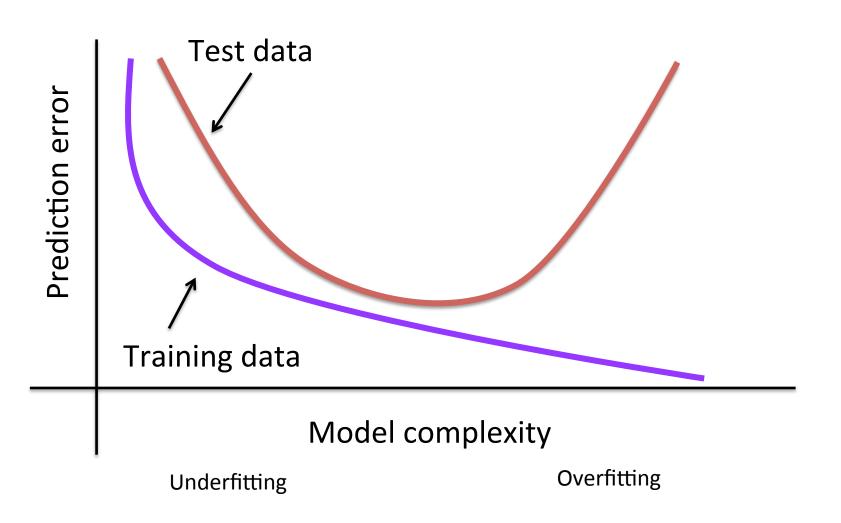
William of Ockham (1285-1349) Principle of Parsimony:

"One should not increase, beyond what is necessary, thenumber of entities required to explain anything."

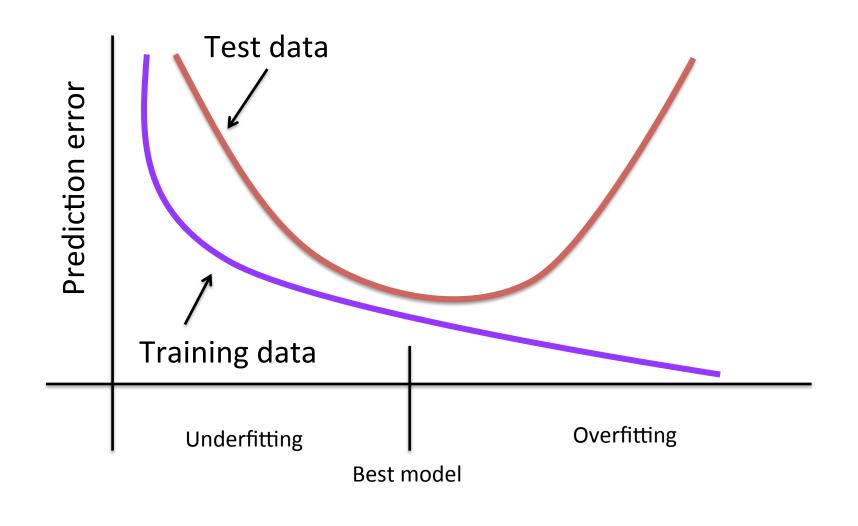


Alternatively: seek the simplest explanation.

## Error and model complexity



# Error and model complexity



# How can we determine the optimal complexity? model selection

- Hold-out method
- Cross-validation
- Complexity regularization
  - Variable selection
  - Variable penalization
- Information criteria (AIC and BIC)

#### Hold-out method

 We would like to pick a model with lowest generalization error.

Simple idea. Find some "validation" data.
 Train model on training data, measure performance of the model on validation data.

Training data

Validation data

$$D_T = \{(\vec{x}_1, y_1), ..., (\vec{x}_n, y_n)\}$$

$$D_V = \{(\vec{x}_1, y_1), ..., (\vec{x}_n, y_n)\}$$

#### Hold-out method

Model that generalizes: low validation error

#### • Problems:

- What if we don't have some validation data put aside?
- Validation error could be misleading, what if we got unlucky on the split of training/validation data?

# Cross-validation (CV)

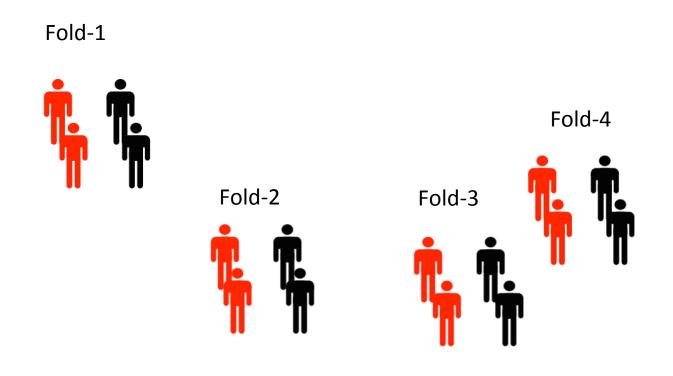
- K-fold cross-validation:
  - 1. Create k partition (folds) of input data
  - 2. Set aside data in one of the partition (fold) for validation
  - 3. Train model on all but the held-out fold
  - 4. Measure cross-validation (CV) error using data from the left-out fold:

$$\mathsf{CV}\;\mathsf{Error}_i = \frac{1}{n_i} \sum_{j \in T_i} L(y_j, \hat{y}_{j(-i)})$$

5. Repeat leaving out for all folds and measure average error

$$\mathsf{CV}\;\mathsf{Error} = \frac{1}{k}\sum_{i=1}^k \mathsf{CV}\;\mathsf{Error}_i$$





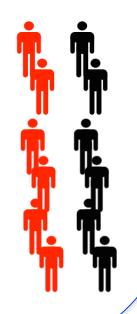
Training data



Validation data







Validation data



Use model to predict class labels for validation fold

Build (train) a model

$$CV\_Error_i = \frac{1}{n_i} \sum_{j \in T_i} L(y_j, \hat{y}_{j(-i)})$$

#### K-fold cross-validation

- k can range from 2 to n:
  - Larger k:
    - Variance of true error will be high
    - Computational time will be large
    - + bias of error estimate will be small
  - Smaller k:
    - + reduced computation time
    - + variance of error estimate is small
    - Bias of error estimate is large
- N-fold CV is called "Leave-One-Out" CV
- In practice:
  - Lower k is more reliable, eg., 3-fold CV is standard.

# How do we measure error on validation/test set?

#### continuous response

#### Examples:

Squared error loss

$$L(\boldsymbol{y}, \hat{\boldsymbol{y}}) = \|\boldsymbol{y} - \hat{\boldsymbol{y}}\|_{2}^{2} = \sum_{i=1}^{n} (y_{i} - \hat{y}_{i})^{2}$$

Absolute error loss

$$L(y, \hat{y}) = \|y - \hat{y}\|_1 = \sum_{i=1}^{n} |y_i - \hat{y}_i|_1$$

# Binary response/labels

- ► True positive (TP)
- True negative (TN)
- False positive (?Type I error?) (FP)
- False negative (?Type II error?) (FN)
- Accuracy (TP + TN)/N (=1-error rate)
- Sensitivity: TP/(TP+FN) (recall)
- Specificity: TN/(TN + FP)

## Concluding remarks

#### Supervised learning:

- Write down a model → objective function
- Algebraically simplify model/objective as much as possible
- Write down the optimization problem: solve for parameters
   (highly recommend doing the above, and program your own logistic regression function)

#### Overfitting:

- Constantly think about it!
- Overfitting can severely delude you about the accuracy of your model

#### Cross-validation:

- Good fist-pass solution for tuning parameters, finding the "right" model
- But could be computationally very expensive
- Serious assumption: test and training data are independent (this may not always be the case, especially due to systematic artifacts in genomics)