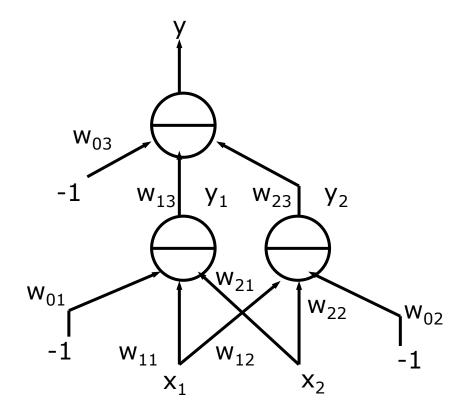
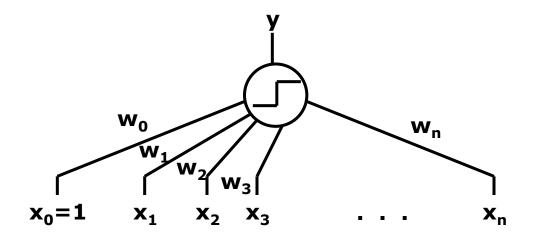
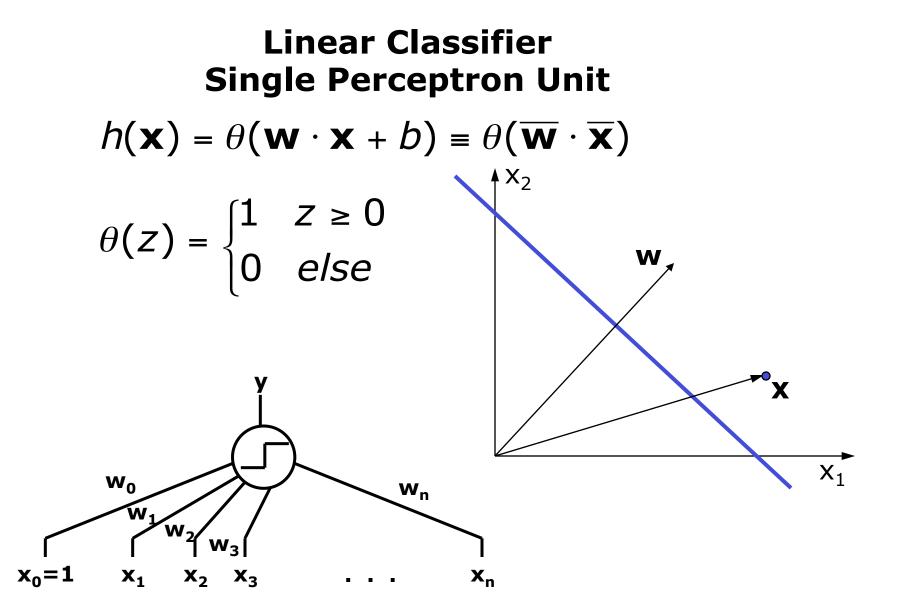
Artificial Neural Networks (Feedforward Nets)

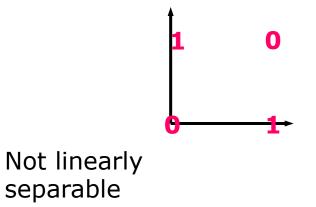


Single Perceptron Unit

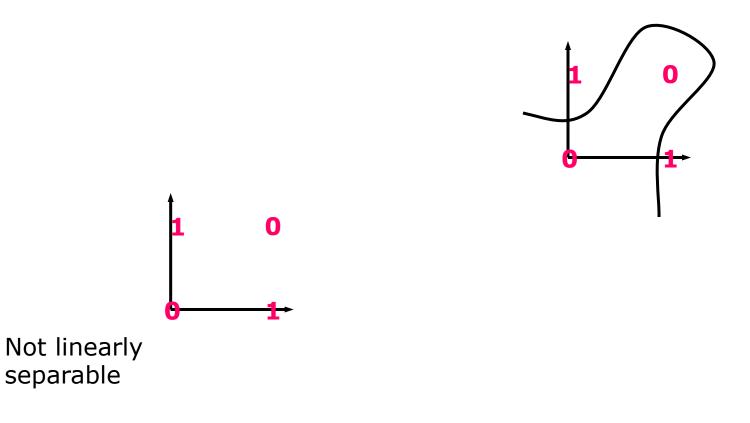




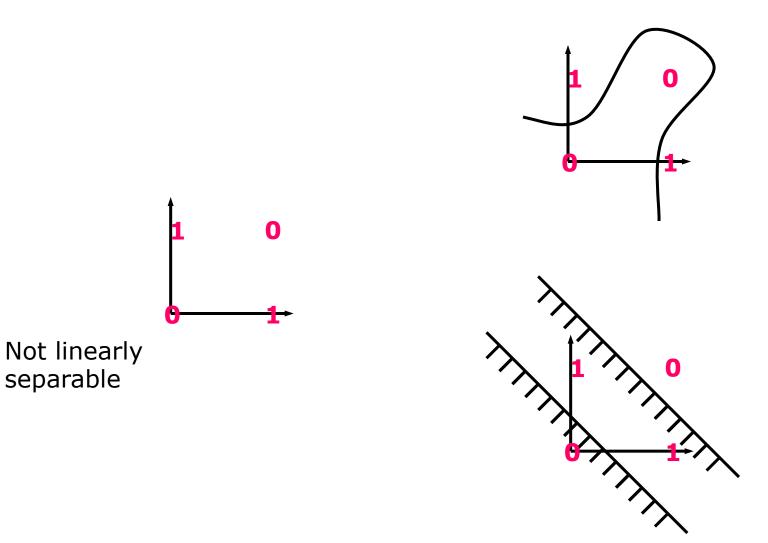
Beyond Linear Separability



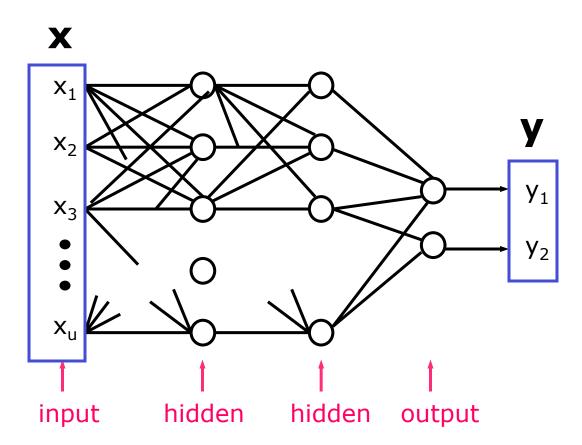
Beyond Linear Separability



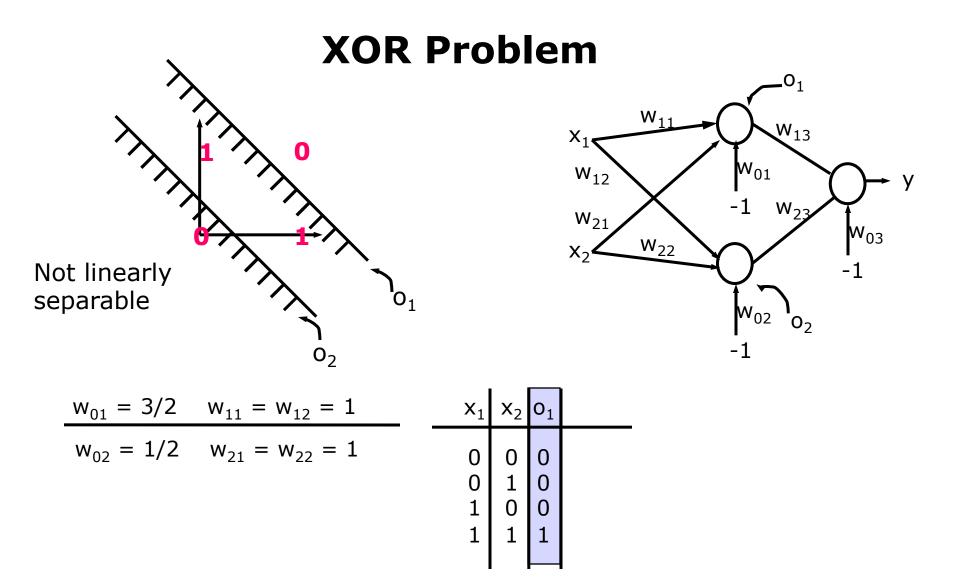
Beyond Linear Separability

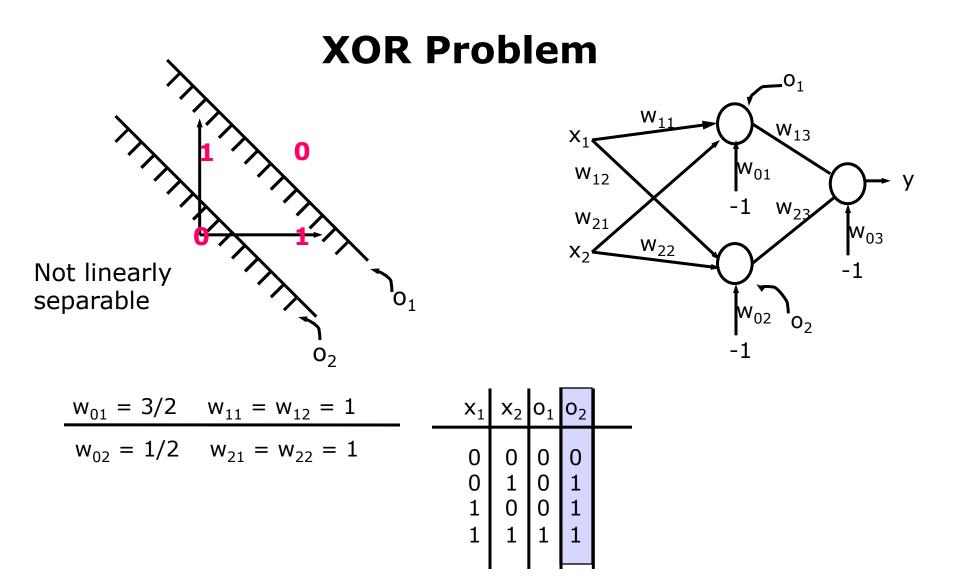


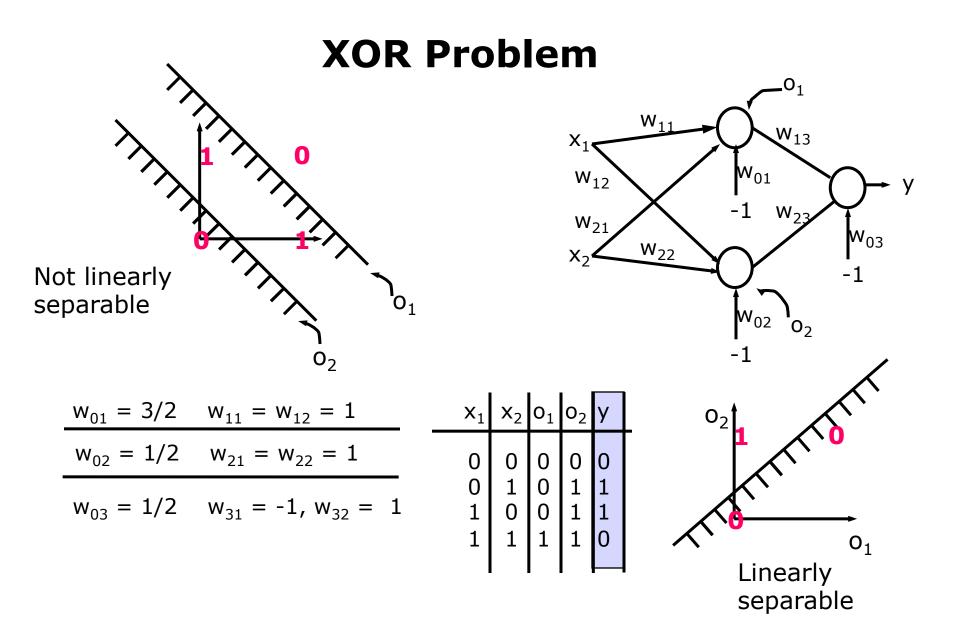
Multi-Layer Perceptron



- More powerful than single layer.
- Lower layers transform the input problem into more tractable (linearly separable) problems for subsequent layers.







- Any set of training points can be separated by a three-layer perceptron network.
- "Almost any" set of points separable by two-layer perceptron network.
- But, no efficient learning rule is known.

- Any set of training points can be separated by a three-layer perceptron network.
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- By no efficient learning rule is known.

May need an exponential number of units.

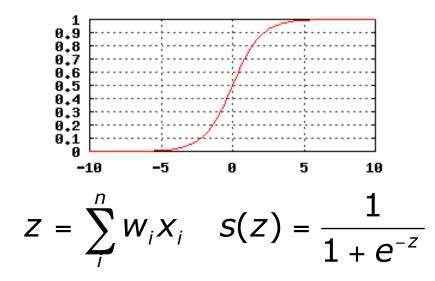
Two "hidden" layers and one output layer

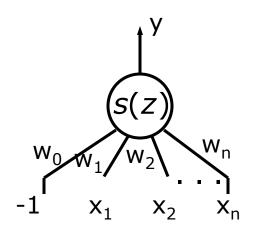
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- Could we use gradient ascent/descent?
- We would need smoothness: small change in weights produces small change in output.
- Threshold function is not smooth.

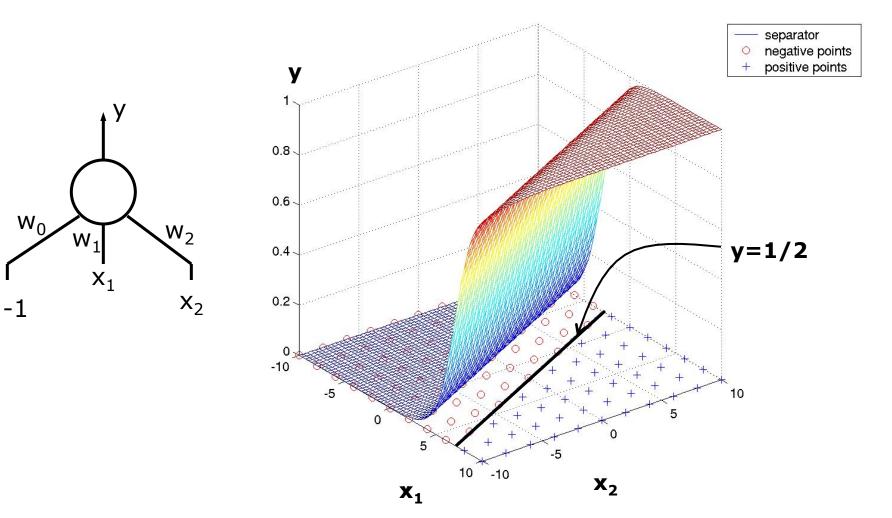
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- Threshold function is not smooth.
- Use a smooth threshold function!

Sigmoid Unit





Sigmoid Unit



Training

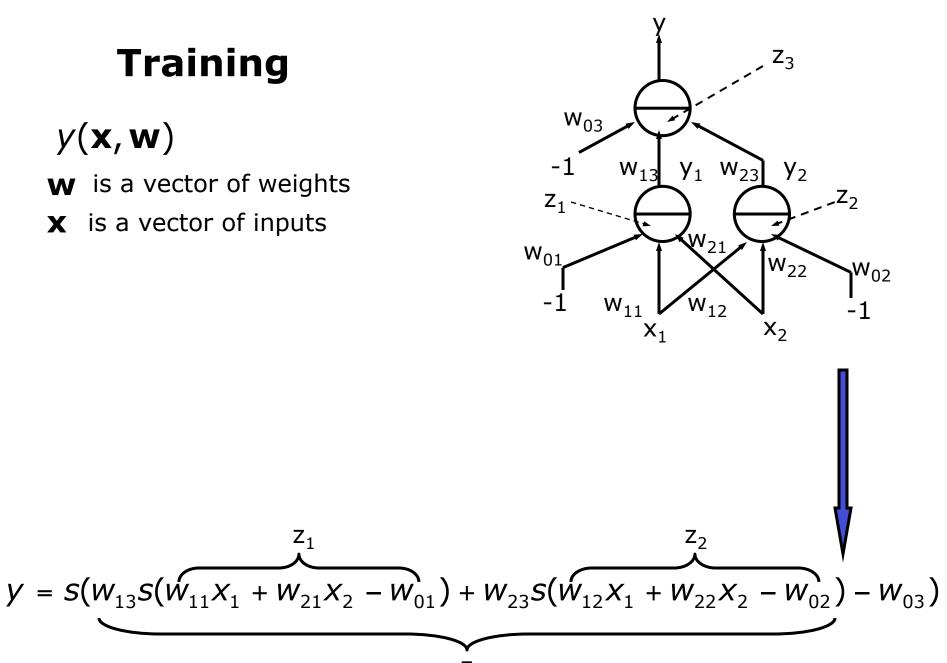
 $y(\mathbf{x}, \mathbf{w})$

w is a vector of weights

 Z_1

 Z_3

X is a vector of inputs



Training

 $y(\mathbf{x}, \mathbf{w})$

w is a vector of weights

 \boldsymbol{X} is a vector of inputs

 \mathbf{Y}' is desired output:

Error over the training set for a given weight vector:

$$E = \frac{1}{2} \sum_{i} (y(\mathbf{x}^{i}, \mathbf{w}) - y^{i})^{2}$$

Our goal is to find weight vector that minimizes error

$$y = S(W_{13}S(W_{11}X_1 + W_{21}X_2 - W_{01}) + W_{23}S(W_{12}X_1 + W_{22}X_2 - W_{02}) - W_{03})$$

W₀₃

Ζı

 W_{01}

Z٦

Y₂

W₂₂

 X_2

 JZ_2

-W₀₂

W₂₃

Y₁

w₂₁

W₁₂

 W_{13}

 W_{11}

 X_1

Gradient Descent

$$E = \frac{1}{2} \sum_{i} (y(\mathbf{x}^{i}, \mathbf{w}) - y^{i})^{2}$$
$$\nabla_{\mathbf{w}} E = \sum_{i} (y(\mathbf{x}^{i}, \mathbf{w}) - y^{i}) \nabla_{\mathbf{w}} y(\mathbf{x}^{i}, \mathbf{w})$$

Error on training set

Gradient of Error

$$\nabla_{\mathbf{w}} \mathbf{y} = \left[\frac{\partial \mathbf{y}}{\partial W_1}, \dots, \frac{\partial \mathbf{y}}{\partial W_n} \right]$$

$$\mathbf{w} \leftarrow \mathbf{w} - \eta \nabla_{\mathbf{w}} E$$

Gradient Descent

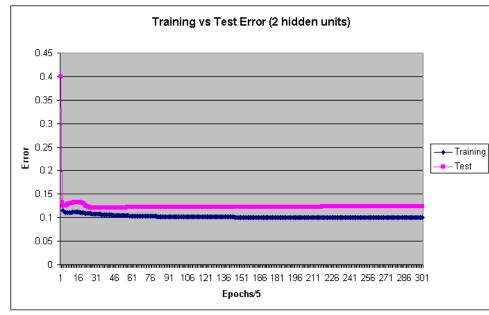
Training Neural Nets without overfitting, hopefully...

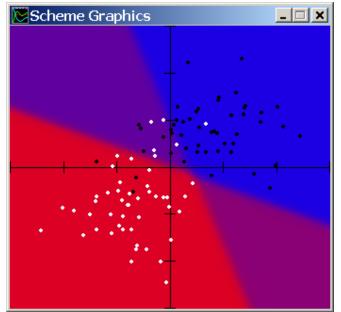
Given: Data set, desired outputs and a neural net with m weights. Find a setting for the weights that will give good predictive performance on new data. Estimate expected performance on new data.

- 1. Split data set (randomly) into three subsets:
 - Training set used for picking weights
 - Validation set used to stop training
 - Test set used to evaluate performance
- 2. Pick random, small weights as initial values
- 3. Perform iterative minimization of error over training set.
- 4. Stop when error on <u>validation set</u> reaches a minimum (to avoid overfitting).
- 5. Repeat training (from step 2) several times (avoid local minima)
- 6. Use best weights to compute error on <u>test set</u>, which is estimate of performance on new data. Do not repeat training to improve this.

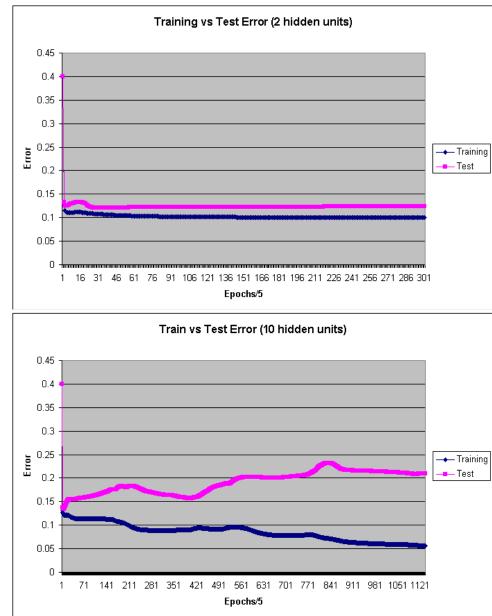
Can use cross-validation if data set is too small to divide into three subsets.

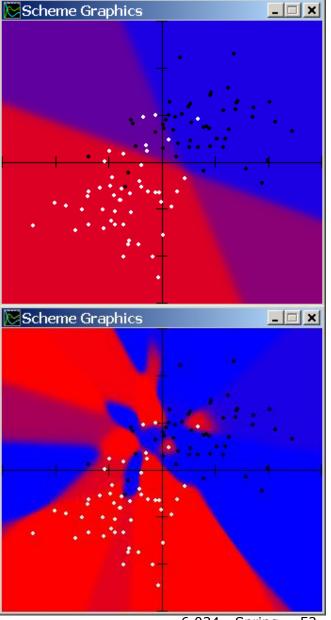
Training vs. Test Error





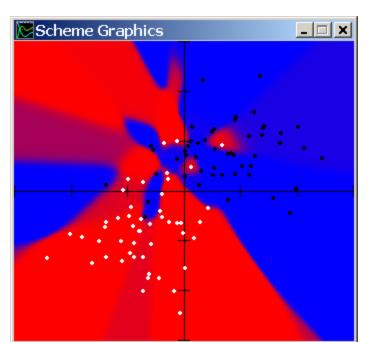
Training vs. Test Error



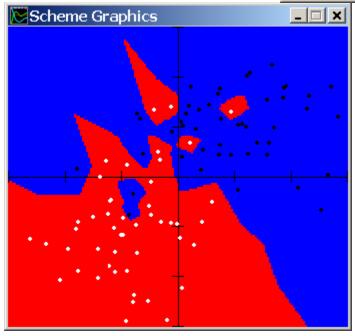


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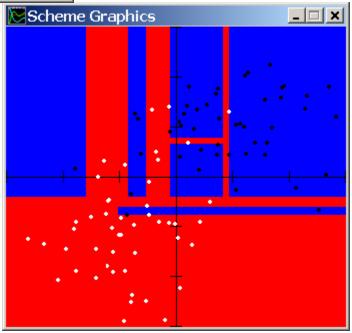
Overfitting is not unique to neural nets...



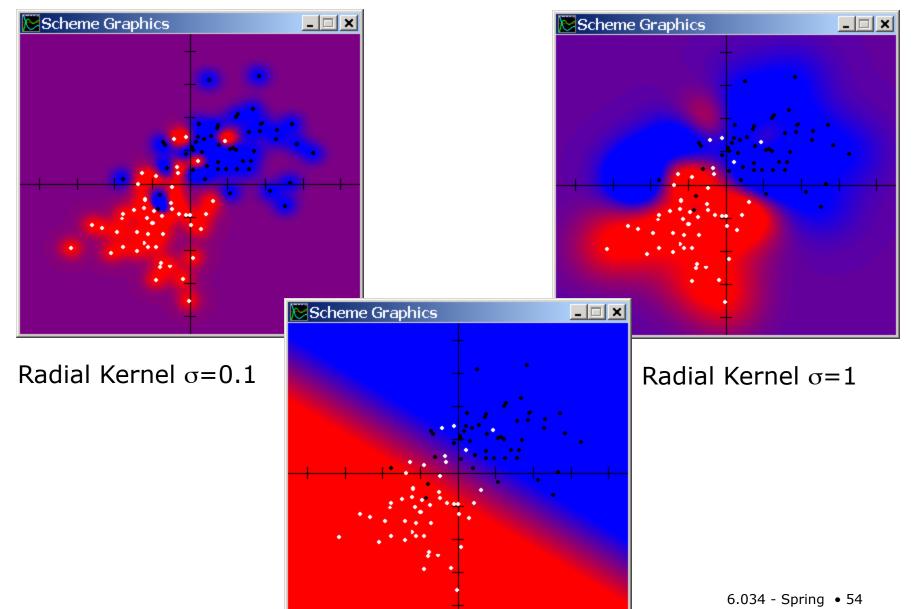
1-Nearest Neighbors



Decision Trees



Overfitting in SVM



On-line vs off-line

There are two approaches to performing the error minimization:

- On-line training present xⁱ and y^{i*} (chosen randomly from the training set). Change the weights to reduce the error on this instance. Repeat.
- Off-line training change weights to reduce the total error on training set (sum over all instances).

On-line training is an approximation to gradient descent since the gradient based on one instance is "noisy" relative to the full gradient (based on all instances). This can be beneficial in pushing the system out of shallow local minima.

Feature Selection

- In many machine learning applications, there are huge numbers of features
 - text classification (# words)
 - gene arrays (5,000 50,000)
 - images (512 x 512 pixels)



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- Too many features
 - make algorithms run slowly
 - risk overfitting
- Find a smaller feature space
 - subset of existing features
 - new features constructed from old ones



Feature Ranking

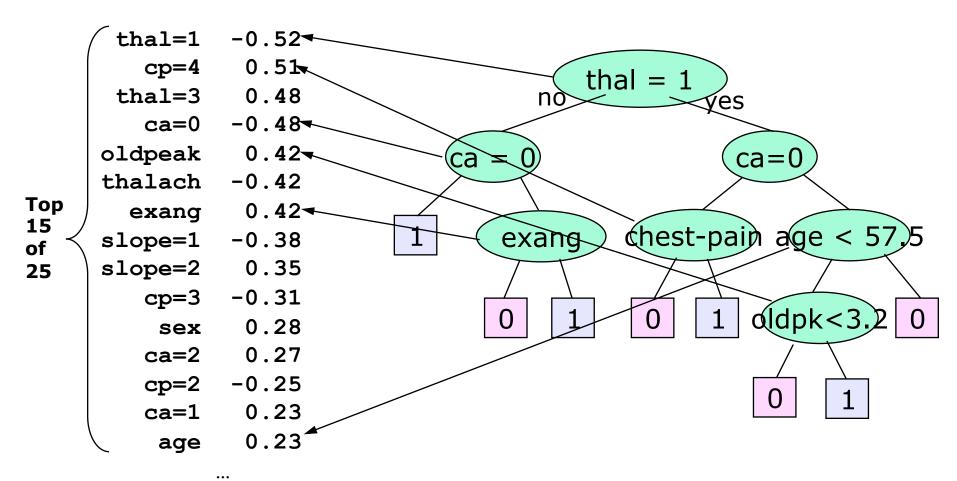
- For each feature, compute a measure of its relevance to the output
- Choose the k features with the highest rankings
- Correlation between feature j and output

$$R(j) = \frac{\sum_{i} (x_{j}^{i} - \overline{x}_{j})(y^{i} - \overline{y})}{\sqrt{\sum_{i} (x_{j}^{i} - \overline{x}_{j})^{2} \sum_{i} (y^{i} - \overline{y})^{2}}}$$
$$\overline{x}_{j} = \frac{1}{n} \sum_{i} x_{j}^{i} \qquad \overline{y} = \frac{1}{n} \sum_{i} y^{i}$$

 Correlation measures how much x tends to deviate from its mean on the same examples on which y deviates from its mean

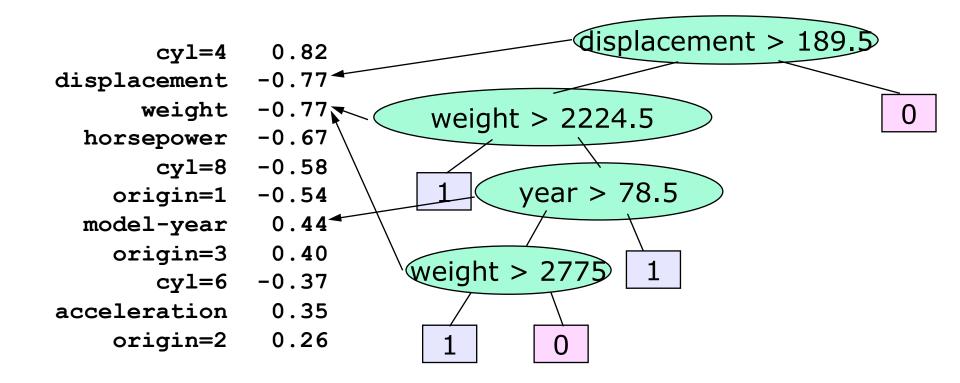


Correlations in Heart Data





Correlations in MPG > 22 data





XOR Bites Back

- As usual, functions with XOR in them will cause us trouble
 - Each feature will, individually, have a correlation of 0 (it occurs positively as much as negatively for positive outputs)
 - To solve XOR, we need to look at groups of features together



Subset Selection

- Consider subsets of variables
 - too hard to consider all possible subsets
 - wrapper methods: use training set or crossvalidation error to measure the goodness of using different feature subsets with your classifier
 - greedily construct a good subset by adding or subtracting features one by one



Forward Selection

Given a particular classifier you want to use
F = {}
For each f_j
Train classifier with inputs F + {f_j}
Add f_j that results in lowest-error classifier
to F
Continue until F is the right size, or error has
quit decreasing



Forward Selection

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• Decision trees, by themselves, do something similar to this



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- Decision trees, by themselves, do something similar to this
- Trouble with XOR



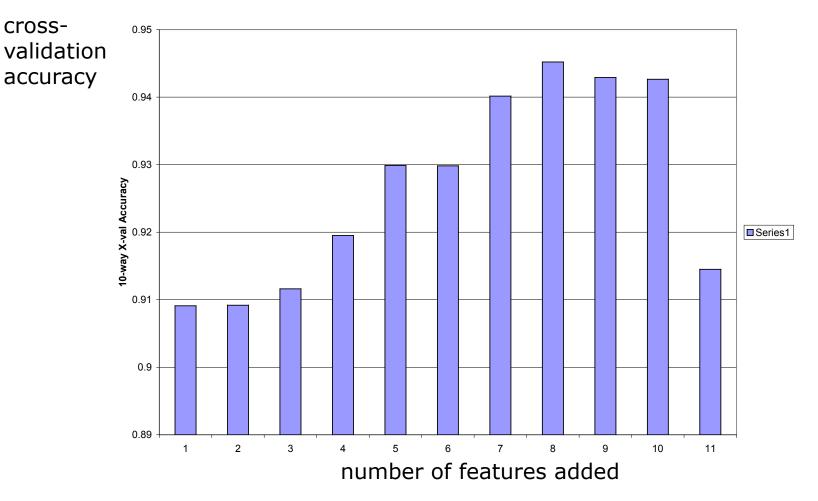
Backward Elimination

Given a particular classifier you want to use
F = all features
For each f;
Train classifier with inputs F - {f;}
Remove f; that results in lowest-error
classifier from F
Continue until F is the right size, or error
increases too much





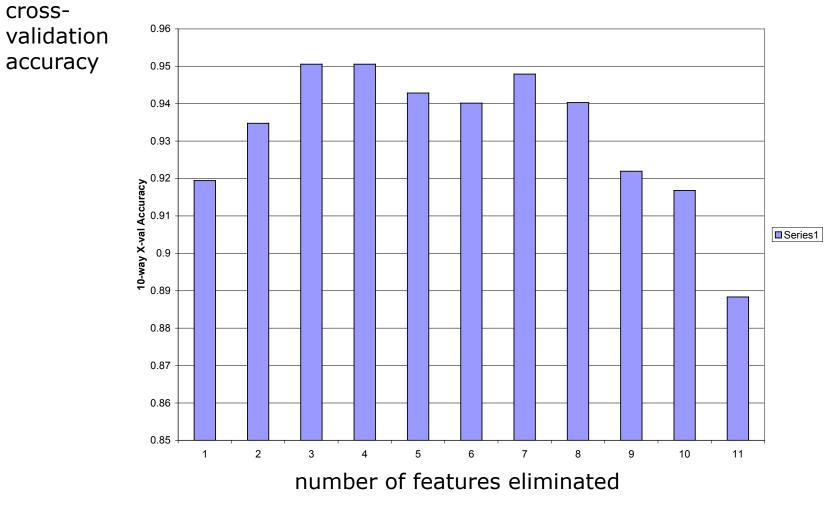
Forward Selection on Auto Data



Forward Selection - Auto



Backward Elimination on Auto Data



Backward Selection - Auto



Forward Selection on Heart Data

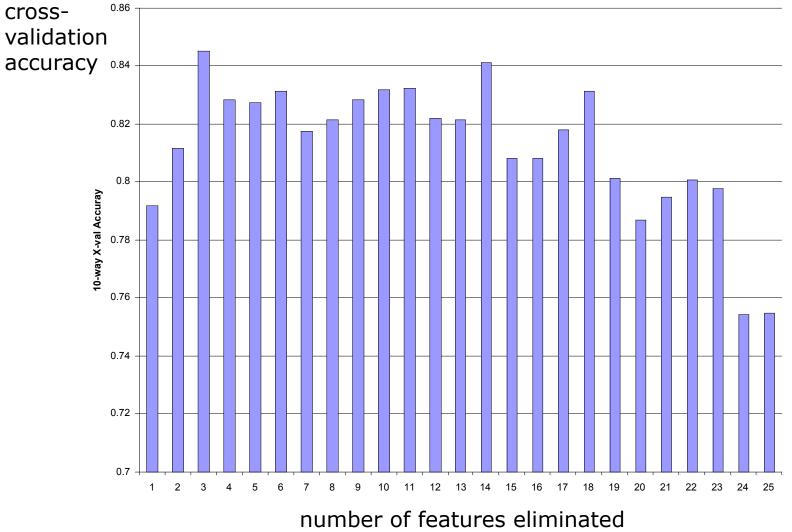
cross-0.86 validation accuracy 0.84 0.82 **10-way X-val Accuracy** 82.0 Series1 0.76 0.74 0.72 10 11 12 13 14 15 16 17 18 19 2 3 4 5 6 7 8 9 20 21 22 23 24 25 1 number of features added

Forward Selection - Hear



Backward Elimination on Heart Data

Backward Elimination - Heart





Recursive Feature Elimination

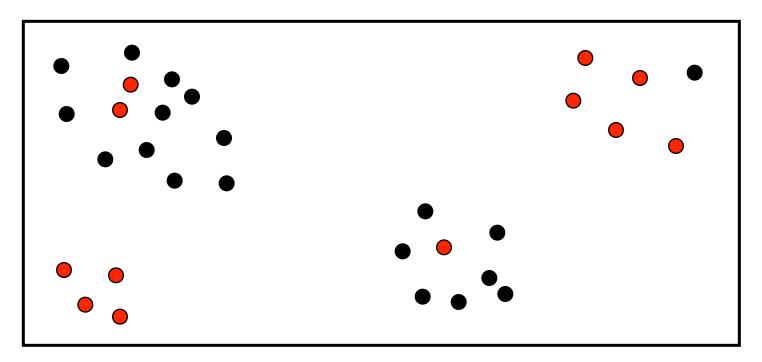
Train a linear SVM or neural network Remove the feature with the smallest weight Repeat

- More efficient than regular backward elimination
- Requires only one training phase per feature



Clustering

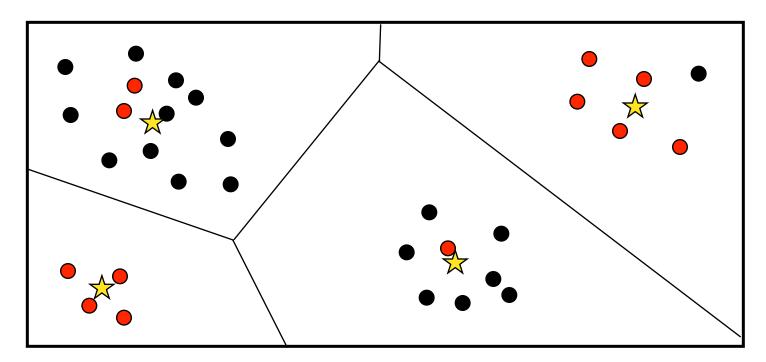
- Form clusters of inputs
- Map the clusters into outputs
- Given a new example, find its cluster, and generate the associated output





Clustering

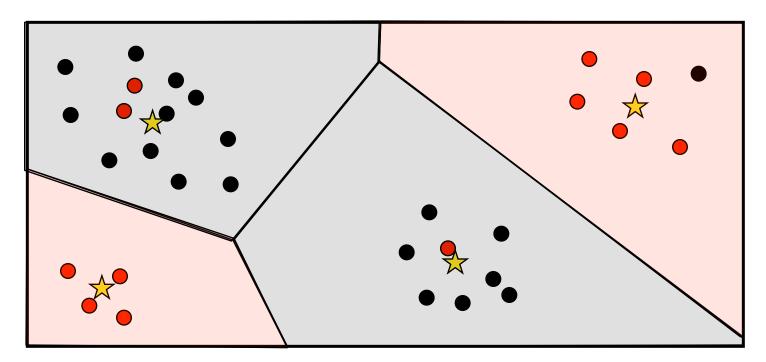
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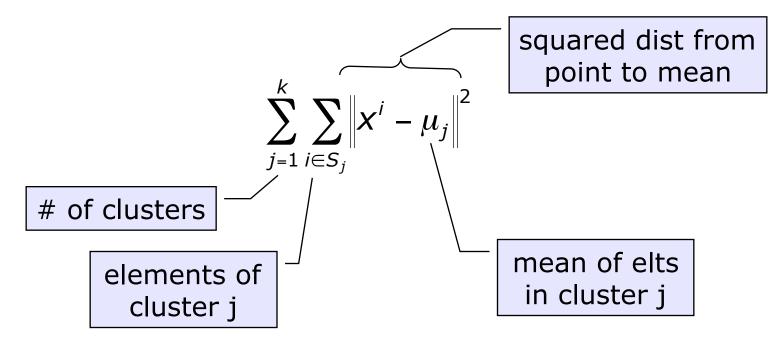
Clustering Criteria

- small distances between points within a cluster
- large distances between clusters
- Need a distance measure, as in nearest neighbor



K-Means Clustering

• Tries to minimize



• Only gets, greedily, to a local optimum



Choose k

Randomly choose k points Cj to be cluster centers



Choose k

Randomly choose k points Cj to be cluster centers Loop

Partition the data into k classes Sj according to which of the Cj they're closest to For each Sj, compute the mean of its elements and let that be the new cluster center



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Guaranteed to terminate



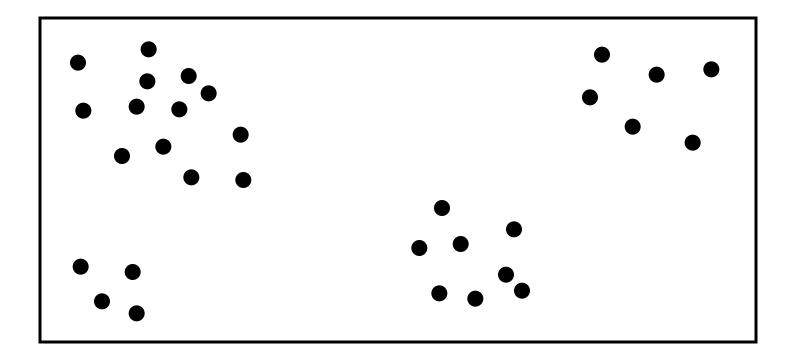
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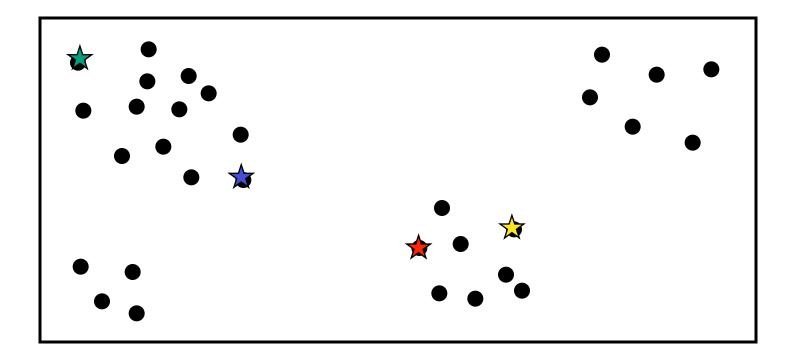
Partition the data into k classes Sj according to which of the Cj they're closest to For each Sj, compute the mean of its elements and let that be the new cluster center Stop when centers quit moving

- Guaranteed to terminate
- If a cluster becomes empty, re-initialize the center

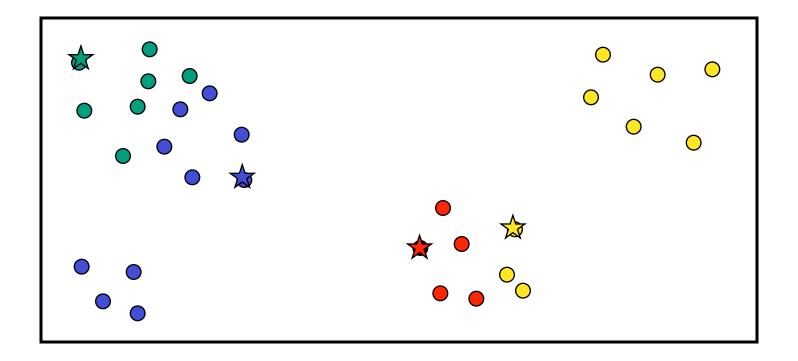




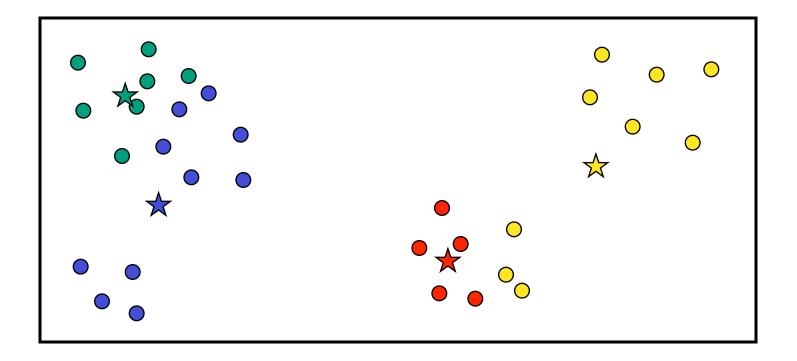




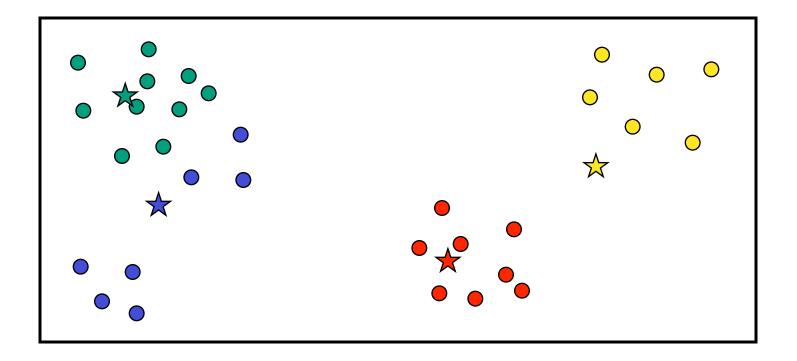




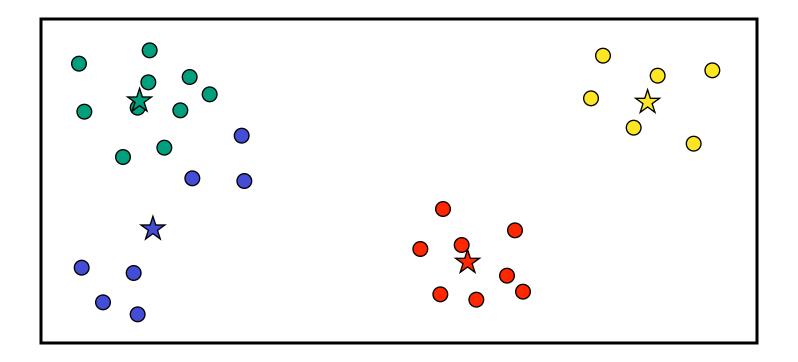




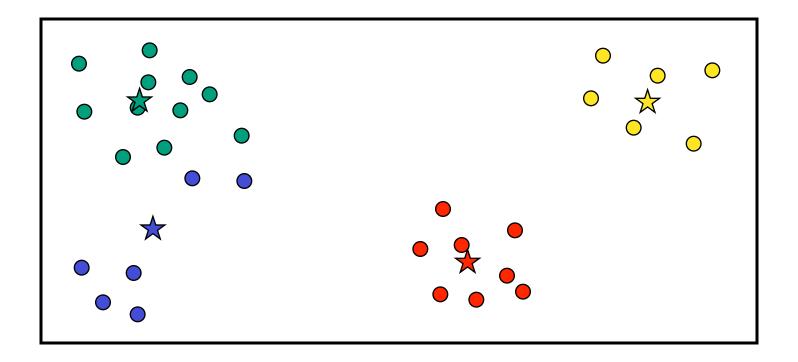




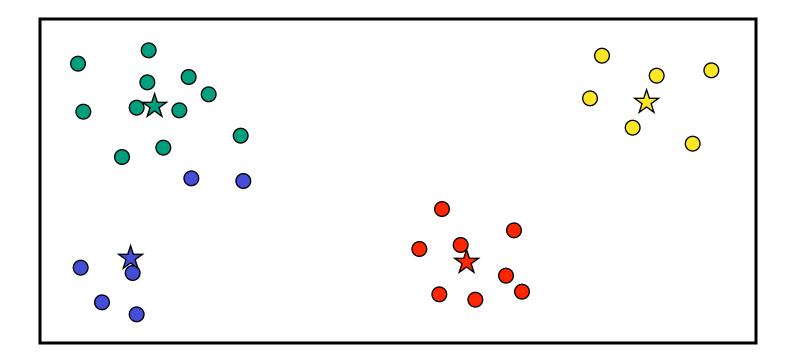




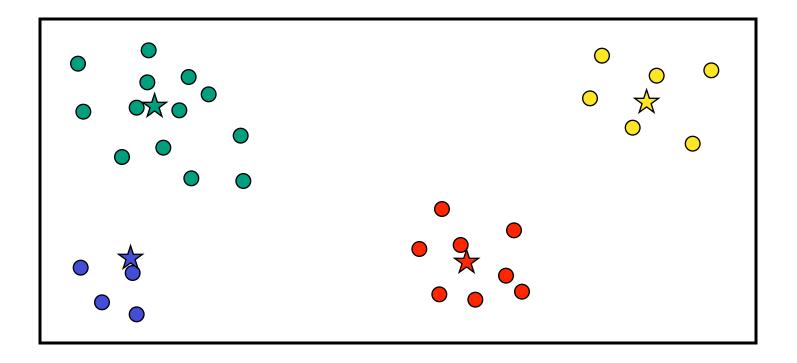




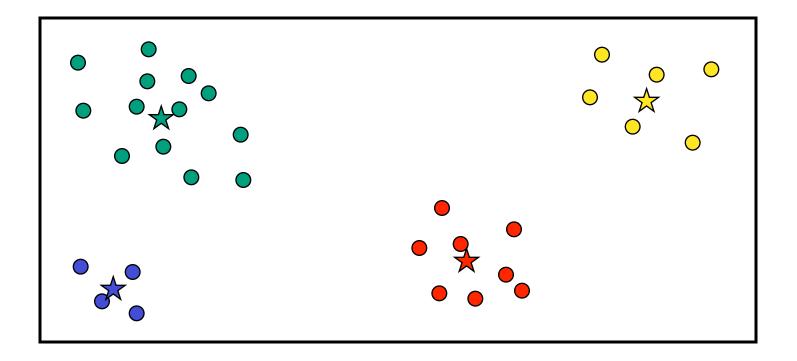














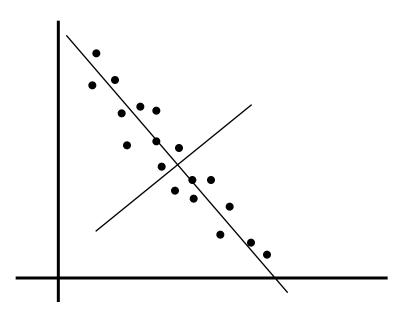
Principal Components Analysis

- Given an n-dimensional real-valued space, data are often nearly restricted to a lower-dimensional subspace
- PCA helps us find such a subspace whose coordinates are linear functions of the originals



Principal Components Analysis

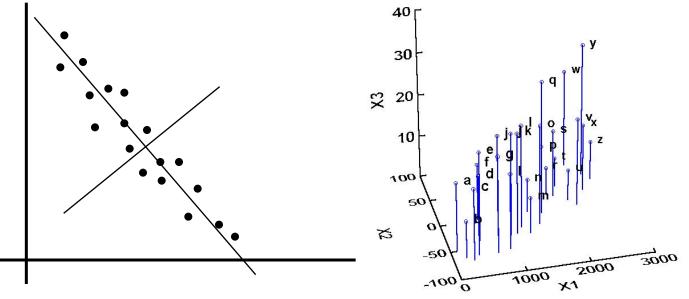
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http://www.okstate.edu/artsci/ botany/ordinate/PCA.htm



Normalize the data (subtract mean, divide by stdev)



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- Find the line along which the data has the most variability: that's the first principal component



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- Repeat



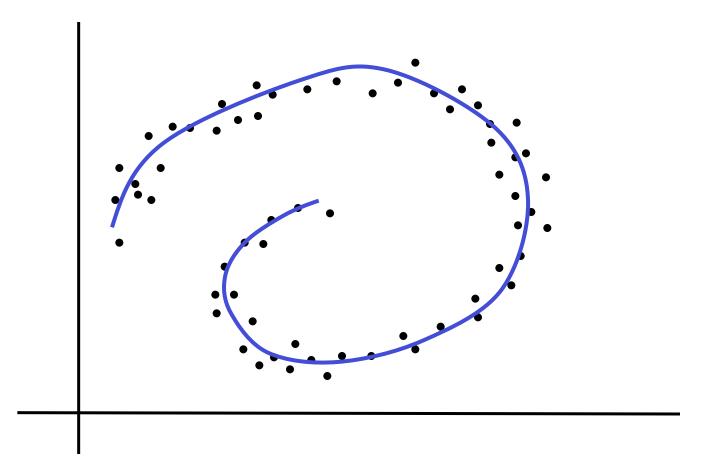
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- Result is a new orthogonal set of axes
- First k give a lower-D space that represents the variability of the data as well as possible



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- Repeat
- Result is a new orthogonal set of axes
- First k give a lower-D space that represents the variability of the data as well as possible
- Really: find the eigenvectors of the covariance matrix with the k largest eigenvalues



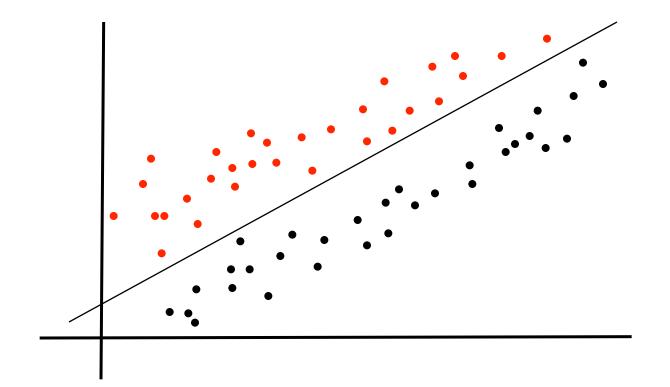
Linear Transformations Only



There are fancier methods that can find this structure

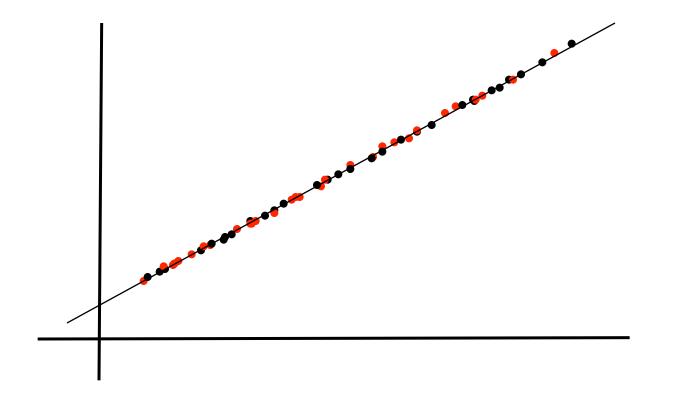


Insensitive to Classification Task



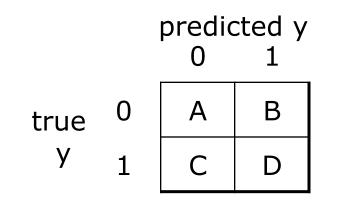


Insensitive to Classification Task

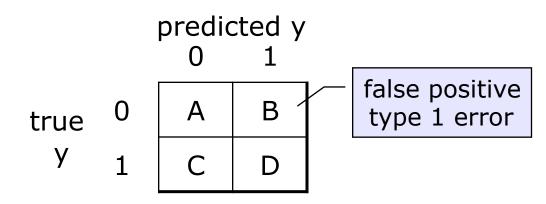


There are fancier methods that can take class into account

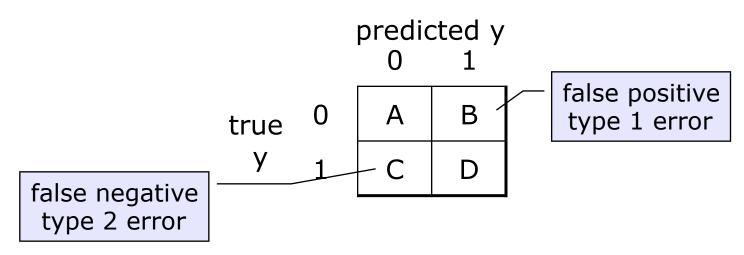




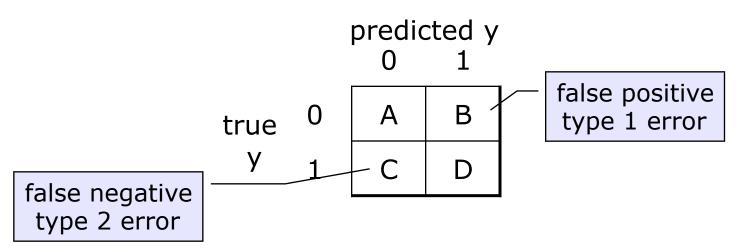






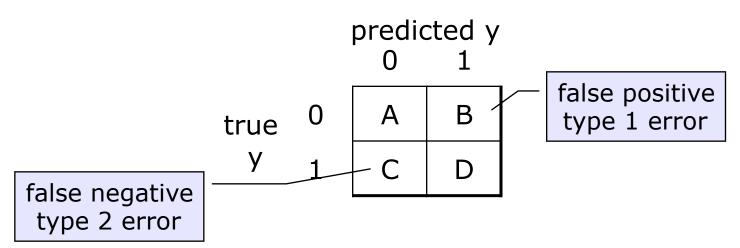






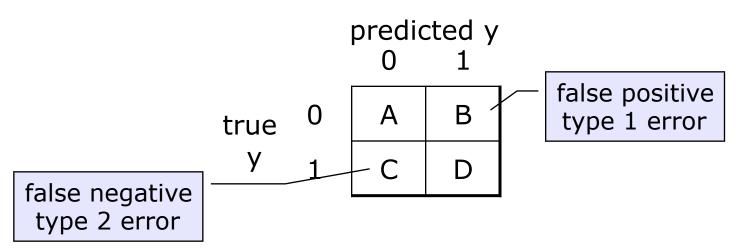
- sensitivity: P(predict 1 | actual 1) = D/(C+D)
 - "true positive rate" (TP)





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- sensitivity: P(predict 1 | actual 1) = D/(C+D)
 - "true positive rate" (TP)
- specificity: P(predict 0 | actual 0) = A/(A+B)
- false-alarm rate: P(predict 1 | actual 0) = B/(A+B)
 - "false positive rate" (FP)



Cost Sensitivity

- Predict whether a patient has pseuditis based on blood tests
 - Disease is often fatal if left untreated
 - Treatment is cheap and side-effect free



Cost Sensitivity

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 - Disease is often fatal if left untreated
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- Which classifier to use?
 - Classifier 1: TP = 0.9, FP = 0.4



Cost Sensitivity

- Predict whether a patient has pseuditis based on blood tests
 - Disease is often fatal if left untreated
 - Treatment is cheap and side-effect free
- Which classifier to use?
 - Classifier 1: TP = 0.9, FP = 0.4
 - Classifier 2: TP = 0.7, FP = 0.1



Build Costs into Classifier

- Assess costs of both types of error
 - use a different splitting criterion for decision trees
 - make error function for neural nets asymmetric; different costs for each kind of error
 - use different values of C for SVMs depending on kind of error



Tunable Classifiers

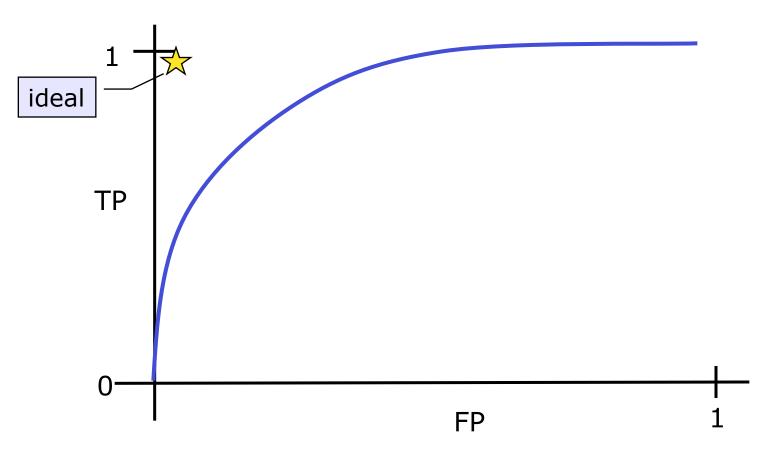
 Classifiers that have a threshold (naïve Bayes, neural nets, SVMs) can be adjusted, post learning, by changing the threshold, to make different tradeoffs between type 1 and type 2 errors



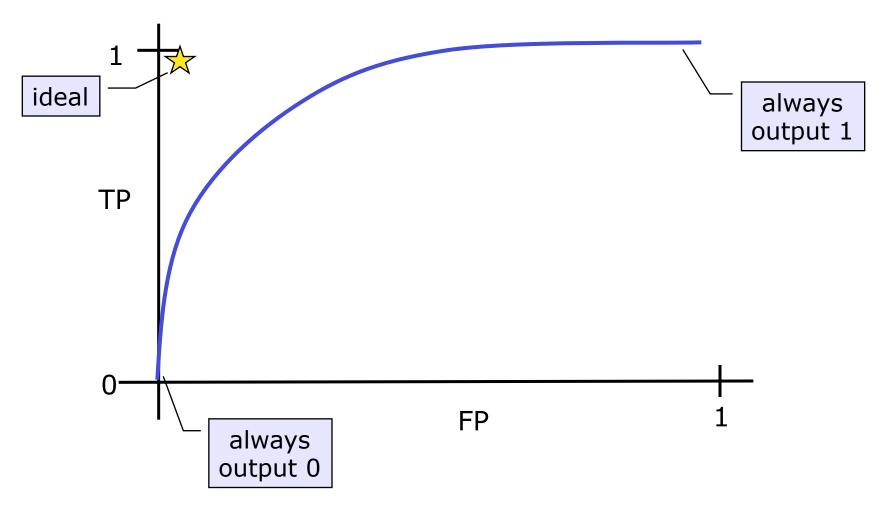
Tunable Classifiers

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 - C₁,C₂: costs of errors
 - P: percentage of positive examples
 - x: tunable threshold
 - TP(x): true positive rate at threshold x
 - FP(x): false positive rate at threshold x
- Expected Cost = $C_2P(1-TP(x)) + C_1(1-P)FP(x)$
- choose x to minimize expected cost

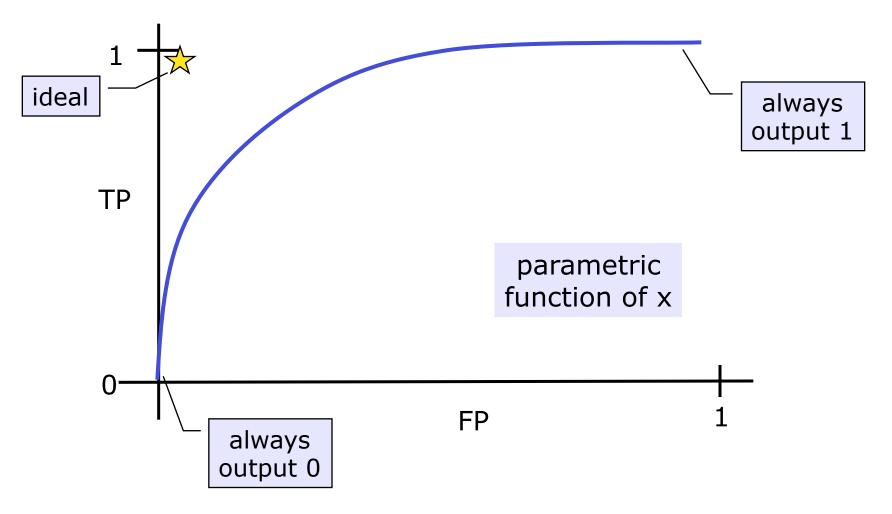




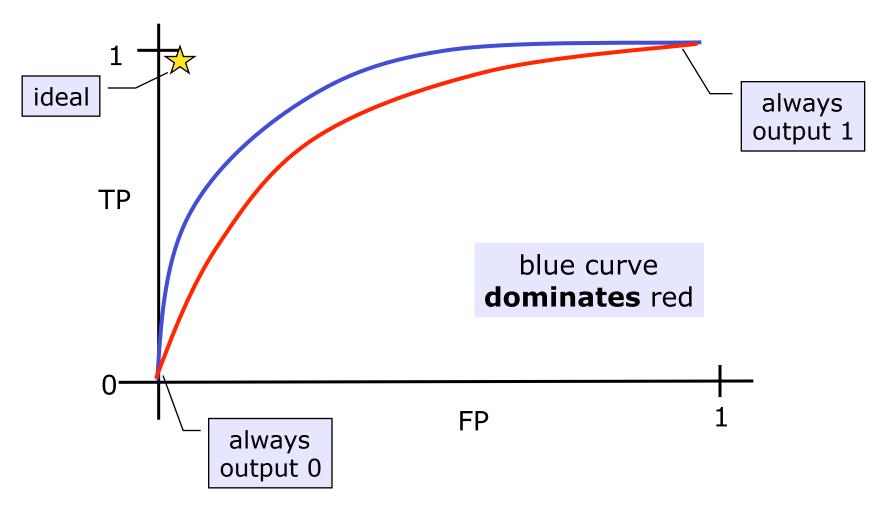














Many more issues!

- Missing data
- Many examples in one class, few in other (fraud detection)
- Expensive data (active learning)



