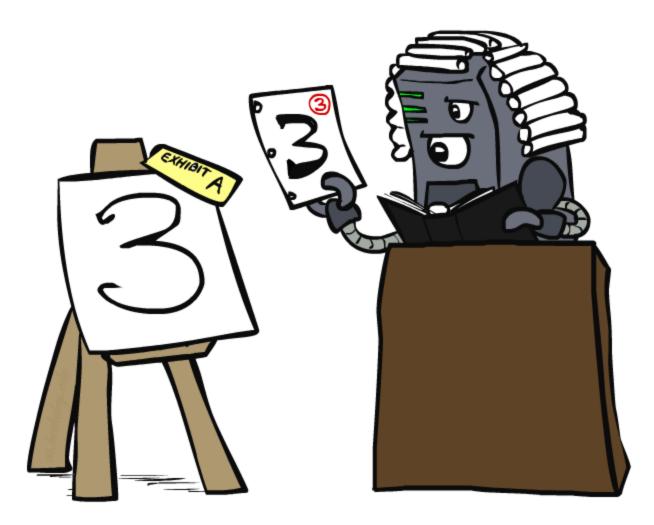
Kernels and Clustering

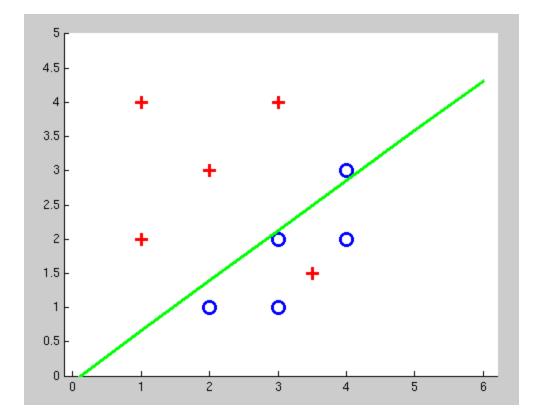
Robert Platt Northeastern University

All slides in this file are adapted from CS188 UC Berkeley

Case-Based Learning



Non-Separable Data



Case-Based Reasoning

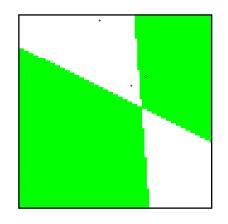
Classification from similarity

- Case-based reasoning
- Predict an instance's label using similar instances
- Nearest-neighbor classification
 - 1-NN: copy the label of the most similar data point
 - K-NN: vote the k nearest neighbors (need a weighting scheme)
 - Key issue: how to define similarity

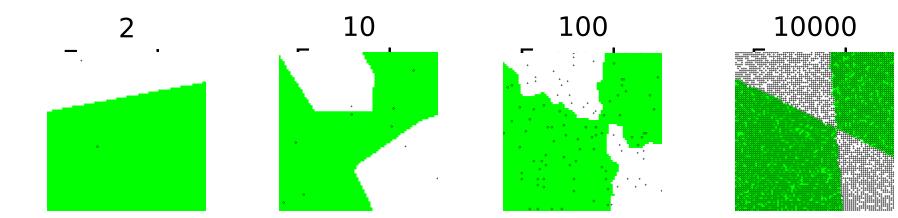


Parametric / Non-Parametric

- Parametric models:
 - Fixed set of parameters
 - More data means better settings
- Non-parametric models:
 - Complexity of the classifier increases with data
 - Better in the limit, often worse in the non-limit
- (K)NN is non-parametric



Truth



Nearest-Neighbor Classification

0

1

2

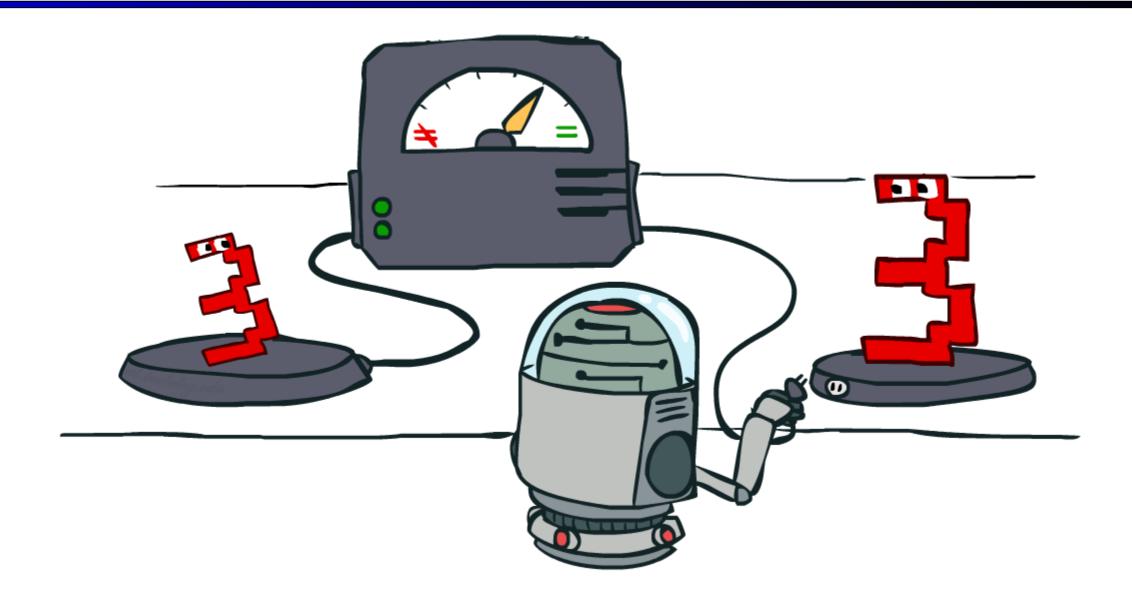
0

- Nearest neighbor for digits:
 - Take new image
 - Compare to all training images
 - Assign based on closest example
- Encoding: image is vector of intensities: $= \langle 0.0 \ 0.0 \ 0.3 \ 0.8 \ 0.7 \ 0.1 \dots 0.0 \rangle$
- What's the similarity function?
 - Dot product of two images vectors?

$$sim(x, x') = x \cdot x' = \sum_{i} x_i x'_i$$

- Usually normalize vectors so ||x|| = 1
- min = 0 (when?), max = 1 (when?)

Similarity Functions



Basic Similarity

Many similarities based on feature dot products:

$$sim(x, x') = f(x) \cdot f(x') = \sum_{i} f_i(x) f_i(x')$$

If features are just the pixels:

$$sim(x, x') = x \cdot x' = \sum_{i} x_i x'_i$$

Note: not all similarities are of this form

Invariant Metrics

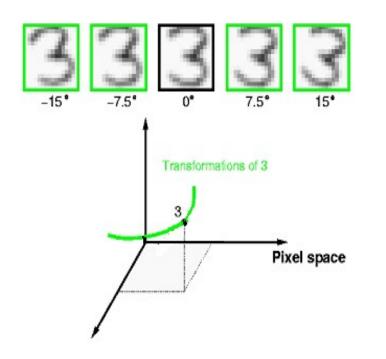
- Better similarity functions use knowledge about vision
- Example: invariant metrics:
 - Similarities are invariant under certain transformations
 - Rotation, scaling, translation, stroke-thickness...
 - E.g:



- 16 x 16 = 256 pixels; a point in 256-dim space
- These points have small similarity in R²⁵⁶ (why?)
- How can we incorporate such invariances into our similarities?

This and next few slides adapted from Xiao Hu,

Rotation Invariant Metrics



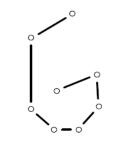
- Each example is now a curve in R²⁵⁶
- Rotation invariant similarity:

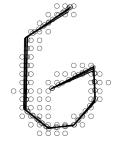
 E.g. highest similarity between images' rotation lines

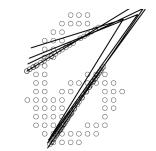
Template Deformation

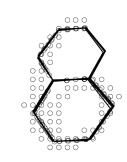
Deformable templates:

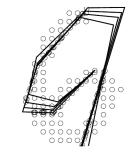
- An "ideal" version of each category
- Best-fit to image using min variance
- Cost for high distortion of template
- Cost for image points being far from distorted terr
- Used in many commercial digit recognizers









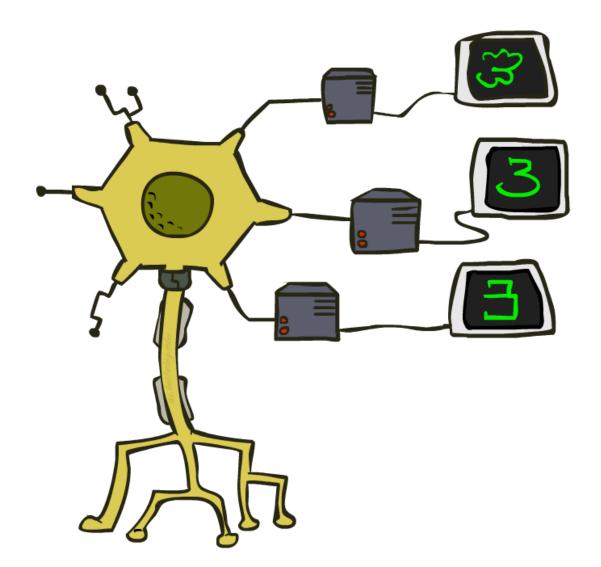


Examples from [Hastie

A Tale of Two Approaches...

- Nearest neighbor-like approaches
 - Can use fancy similarity functions
 - Don't actually get to do explicit learning
- Perceptron-like approaches
 - Explicit training to reduce empirical error
 - Can't use fancy similarity, only linear
 - Or can they? Let's find out!

Kernelization



Perceptron Weights

- What is the final value of a weight w, of a perceptron?
 - Can it be any real vector?
 - No! It's built by adding up inputs.

$$w_y = \mathbf{0} + f(x_1) - f(x_5) + \dots$$

$$w_y = \sum_i \alpha_{i,y} f(x_i)$$

• Can reconstruct weight vectors (the primal representation) from update counts (the dual representation) $\alpha_y = \langle \alpha_{1,y} \ \alpha_{2,y} \ \dots \ \alpha_{n,y} \rangle$

Dual Perceptron

How to classify a new example x?

score
$$(y, x)$$
 = $w_y \cdot f(x)$
= $\left(\sum_i \alpha_{i,y} f(x_i)\right) \cdot f(x)$
= $\sum_i \alpha_{i,y} (f(x_i) \cdot f(x))$
= $\sum_i \alpha_{i,y} K(x_i, x)$

If someone tells us the value of K for each pair of examples, never need to build the weight vectors (or the feature vectors)!

Dual Perceptron

- Start with zero counts (alpha)
- Pick up training instances one by one
- Try to classify x_n ,

$$y = \arg\max_{y} \sum_{i} \alpha_{i,y} K(x_i, x_n)$$

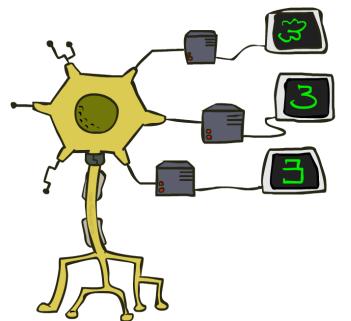
- If correct, no change!
- If wrong: lower count of wrong class (for this instance), raise count of right class (for this instance)

$$\alpha_{y,n} = \alpha_{y,n} - 1 \qquad \qquad w_y = w_y - f(x_n) \alpha_{y^*,n} = \alpha_{y^*,n} + 1 \qquad \qquad w_{y^*} = w_{y^*} + f(x_n)$$

Kernelized Perceptron

- If we had a black box (kernel) K that told us the dot product of two examples x and x':
 - Could work entirely with the dual representation
 - No need to ever take dot products ("kernel trick") SCOPE $(y, x) = w_y \cdot f(x)$

$$= \sum_{i} \alpha_{i,y} K(x_i, x)$$



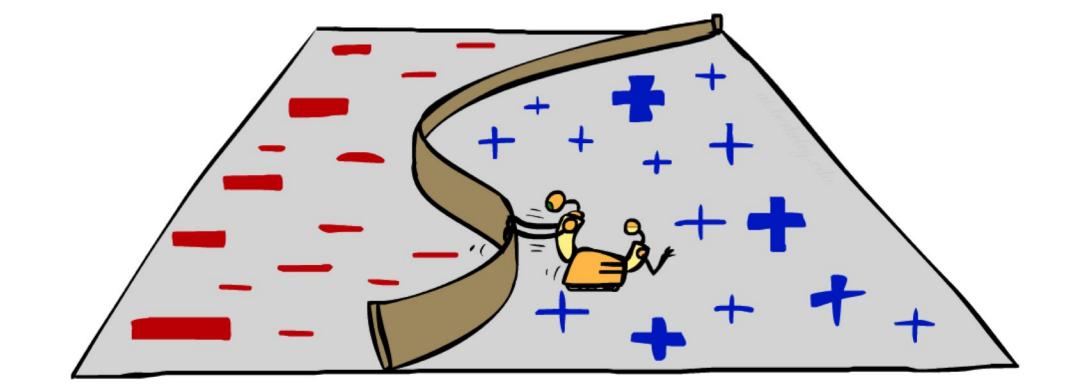
- Like nearest neighbor work with black-box similarities
- Downside: slow if many examples get nonzero alpha

Kernels: Who Cares?

- So far: a very strange way of doing a very simple calculation
- "Kernel trick": we can substitute any* similarity function in place of the dot product
- Lets us learn new kinds of hypotheses

* Fine print: if your kernel doesn't satisfy certain technical requirements, lots of proofs break. E.g. convergence, mistake bounds. In practice, illegal kernels *sometimes* work (but not always).

Non-Linearity

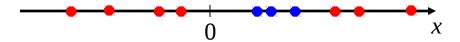


Non-Linear Separators

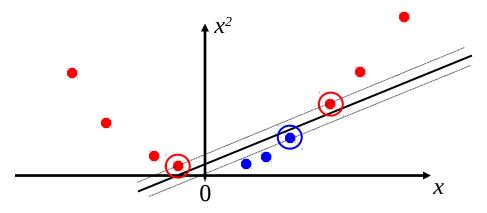
 Data that is linearly separable works out great for linear decision rules:

X

But what are we going to do if the dataset is just too hard?



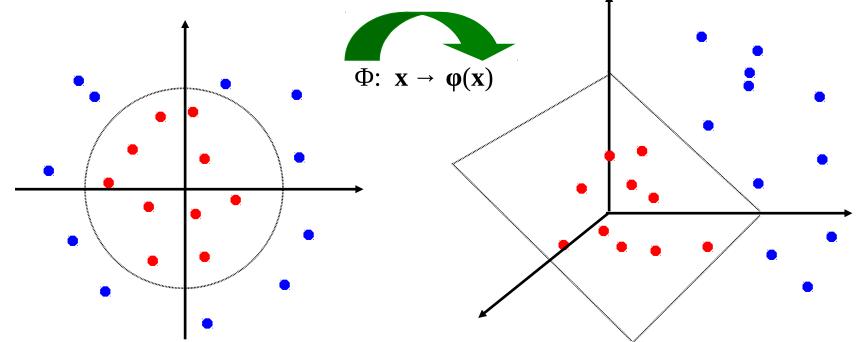
How about... mapping data to a higher-dimensional space:



This and next few slides adapted from Ray Mooney, UT

Non-Linear Separators

General idea: the original feature space can always be mapped to some higher-dimensional feature space where the training set is separable:



Some Kernels

 Kernels implicitly map original vectors to higher dimensional spaces, take the dot product there, and hand the result back

• Linear kernel:
$$K(x, x') = x' \cdot x' = \sum_{i} x_i x'_i$$

• Quadratic kernel: $K(x, x') = (x \cdot x' + 1)^2$

$$= \sum_{i,j} x_i x_j x_i' x_j' + 2 \sum_i x_i x_i' + 1$$

RBF: infinite dimensional representation

$$K(x, x') = \exp(-||x - x'||^2)$$

Discrete kernels: e.g. string kernels

Why Kernels?

- Can't you just add these features on your own (e.g. add all pairs of features instead of using the quadratic kernel)?
 - Yes, in principle, just compute them
 - No need to modify any algorithms
 - But, number of features can get large (or infinite)
 - Some kernels not as usefully thought of in their expanded representation, e.g. RBF kernels
- Kernels let us compute with these features implicitly
 - Example: implicit dot product in quadratic kernel takes much less space and time per dot product
 - Of course, there's the cost for using the pure dual algorithms: you need to compute the similarity to every training datum

Recap: Classification

- Classification systems:
 - Supervised learning
 - Make a prediction given evidence
 - We've seen several methods for this
 - Useful when you have labeled data

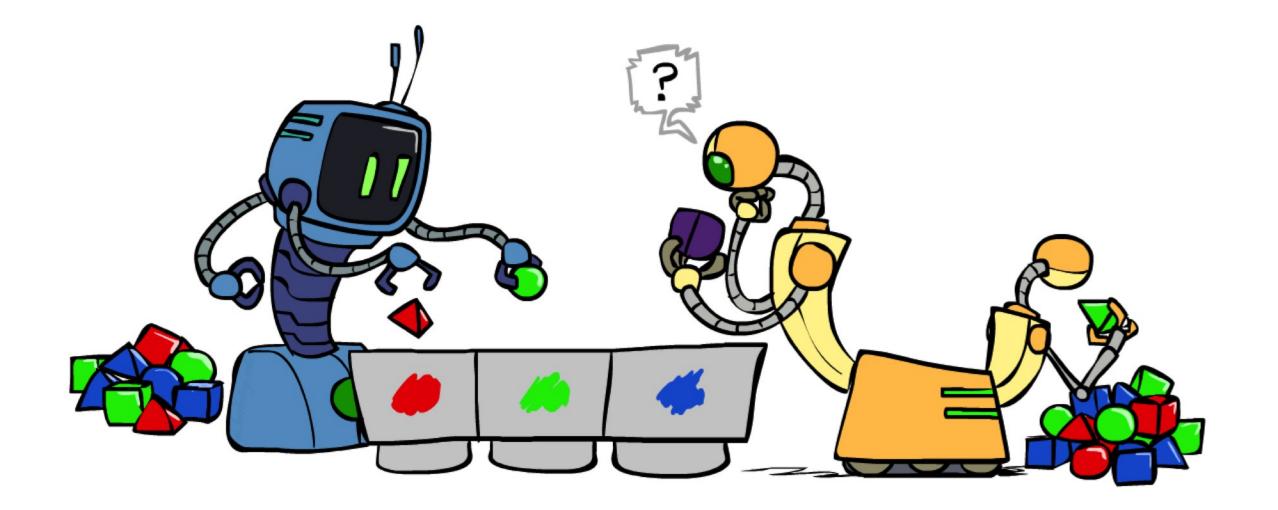


Clustering

- Clustering systems:
 - Unsupervised learning
 - Detect patterns in unlabeled data
 - E.g. group emails or search results
 - E.g. find categories of customers
 - E.g. detect anomalous program executions
 - Useful when don't know what you're looking for
 - Requires data, but no labels
 - Often get gibberish

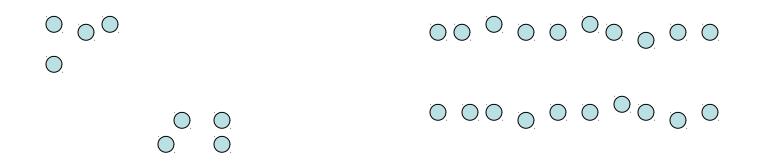


Clustering



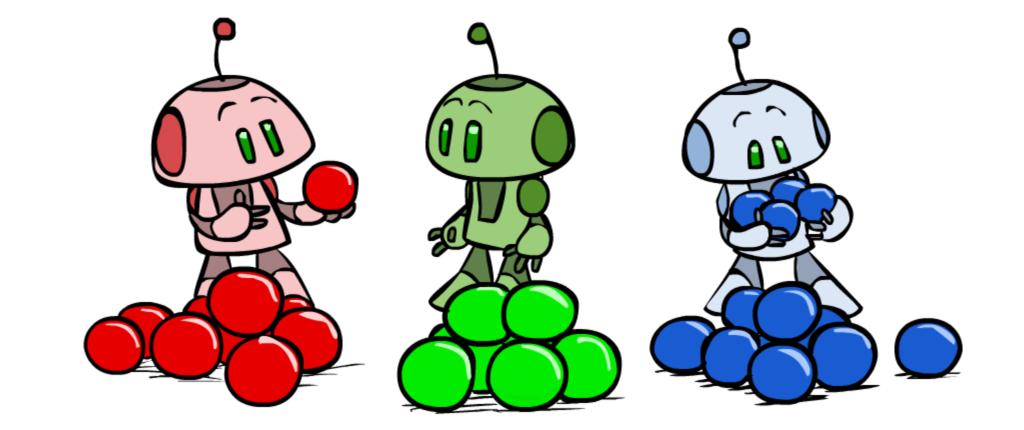
Clustering

- Basic idea: group together similar instances
- Example: 2D point patterns

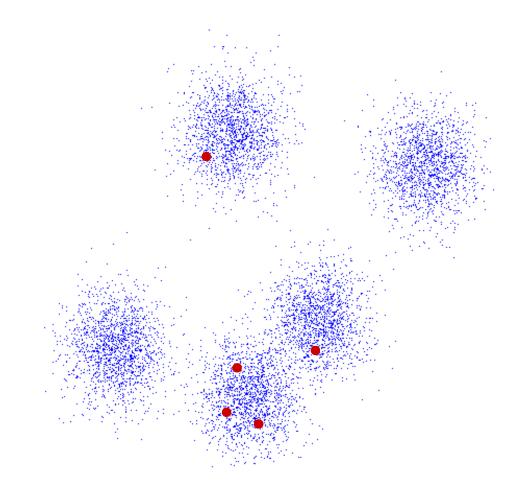


- What could "similar" mean?
 - One option: small (squared) Euclidean distance

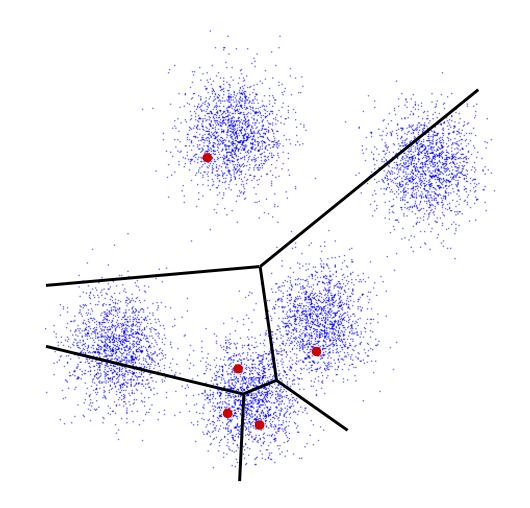
dist
$$(x, y) = (x - y)^{\top} (x - y) = \sum_{i} (x_i - y_i)^2$$



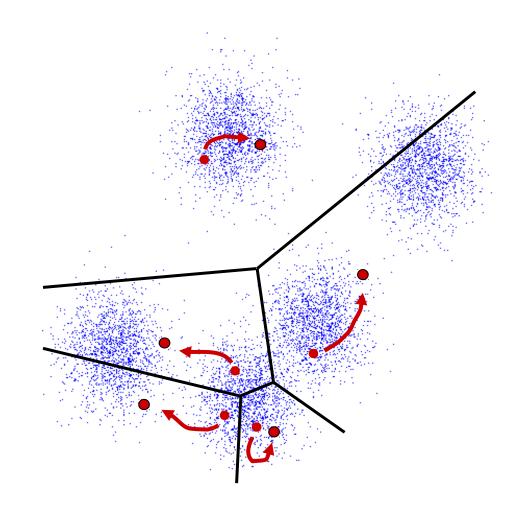
- An iterative clustering algorithm
 - Pick K random points as cluster centers (means)
 - Alternate:
 - Assign data instances to closest mean
 - Assign each mean to the average of its assigned points
 - Stop when no points' assignments change

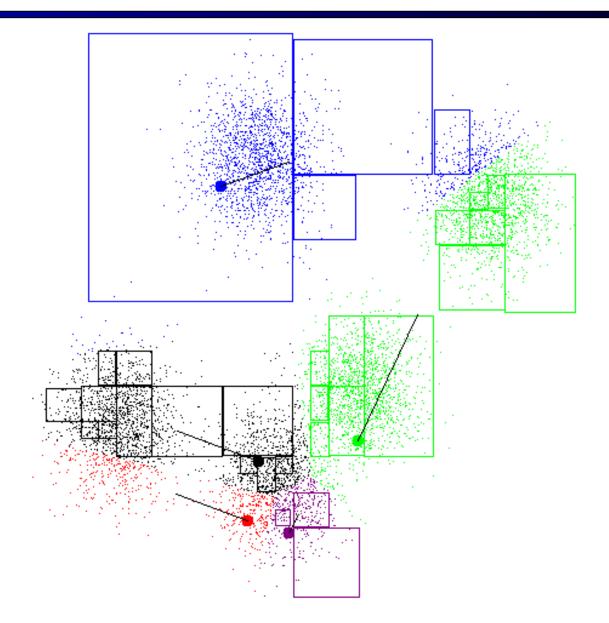


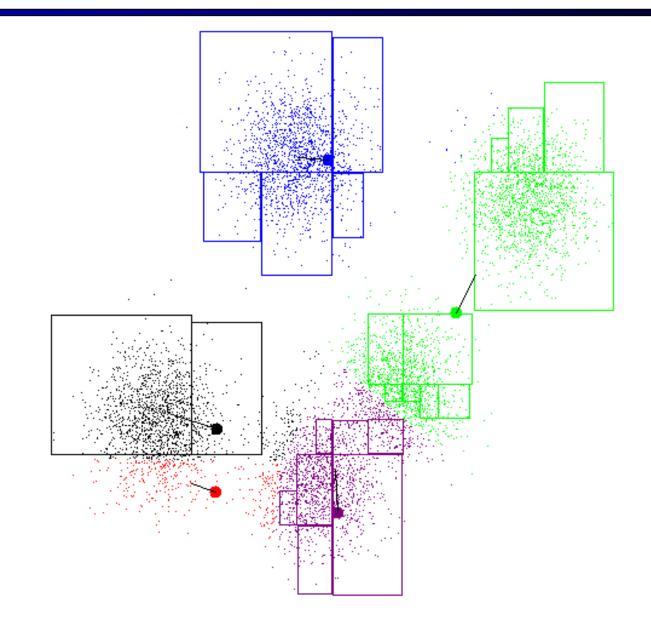
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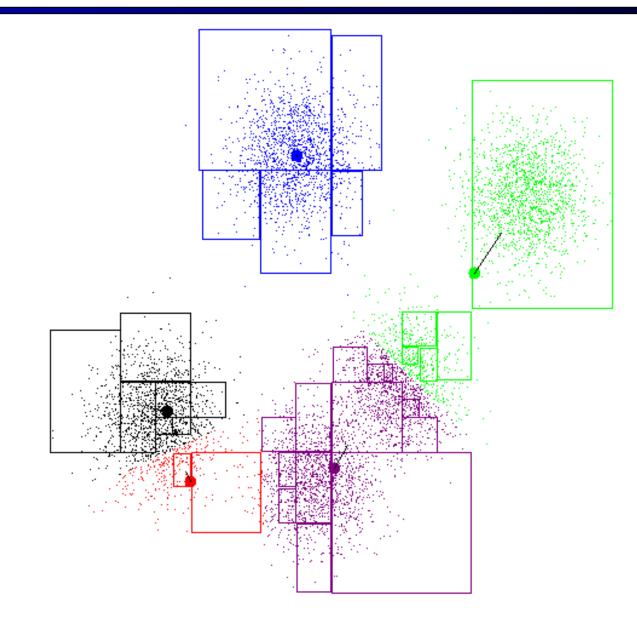


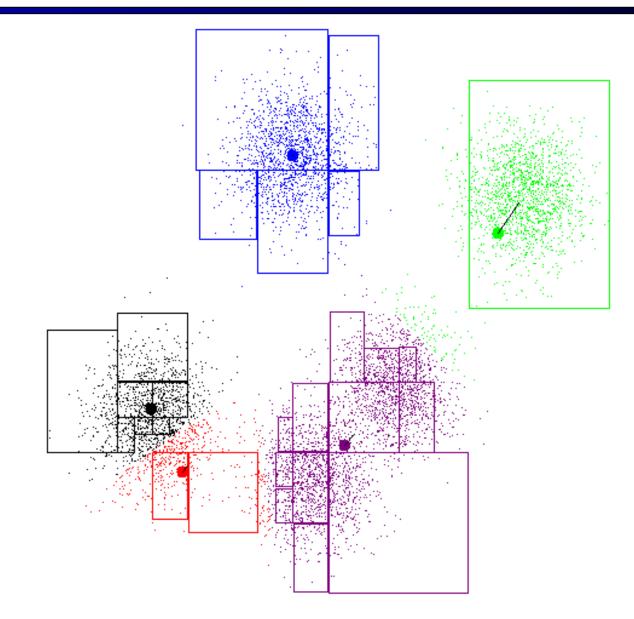
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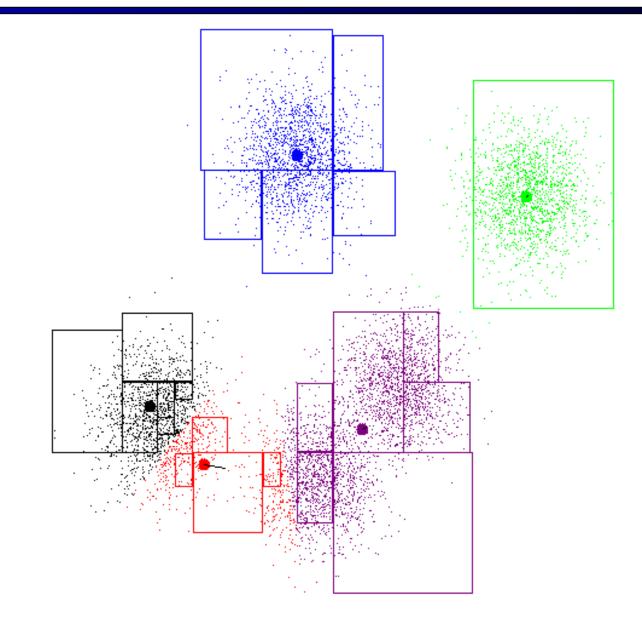


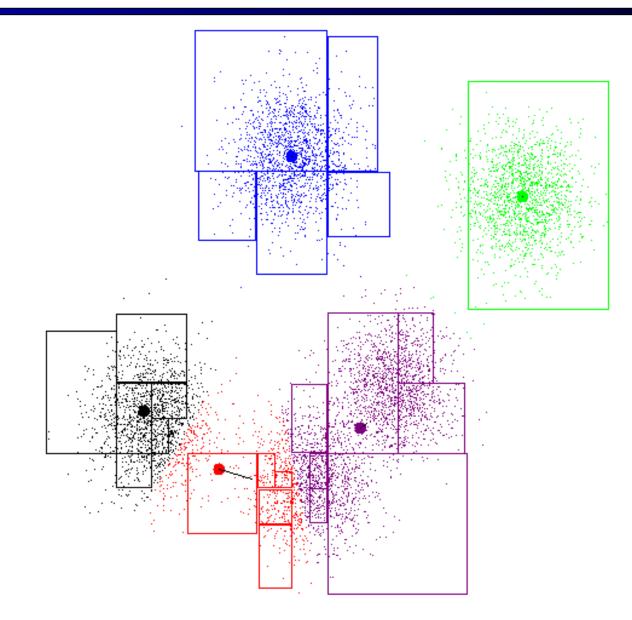


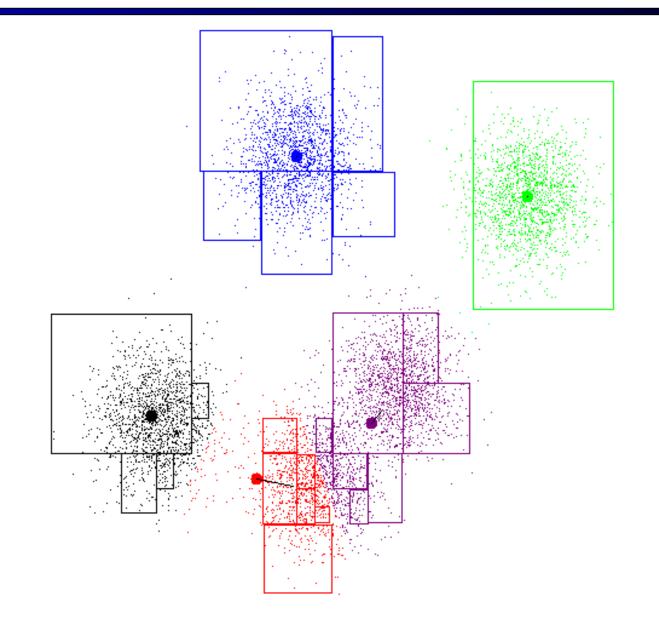


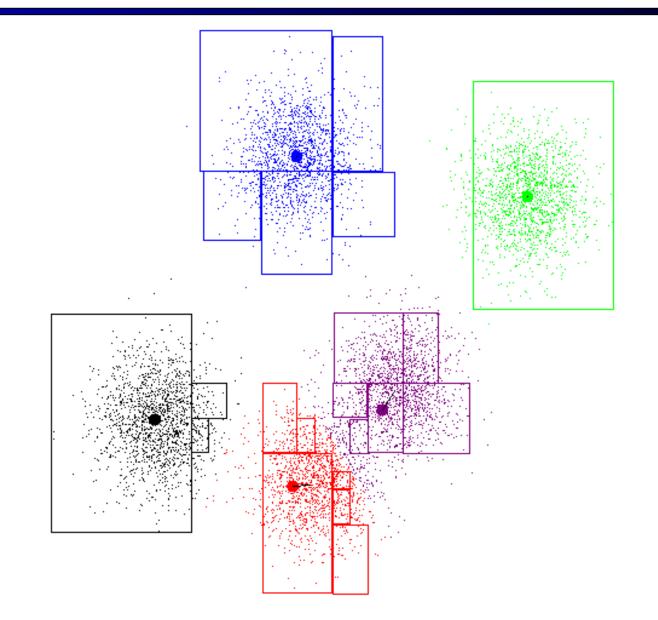


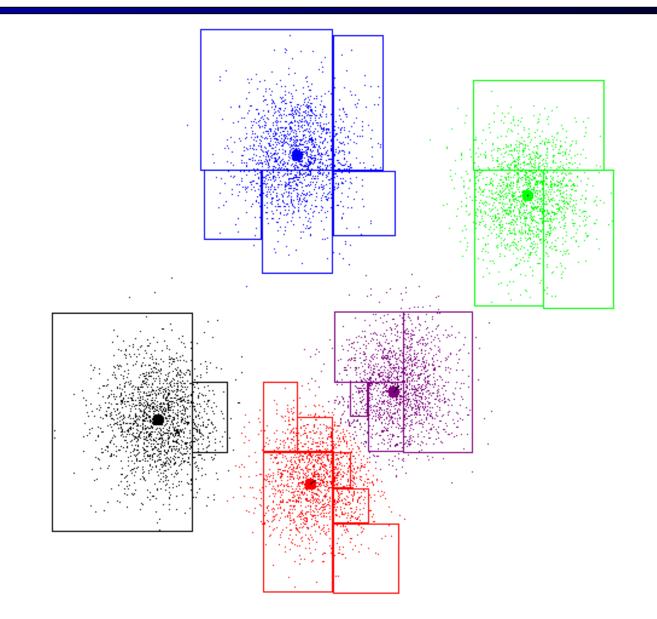










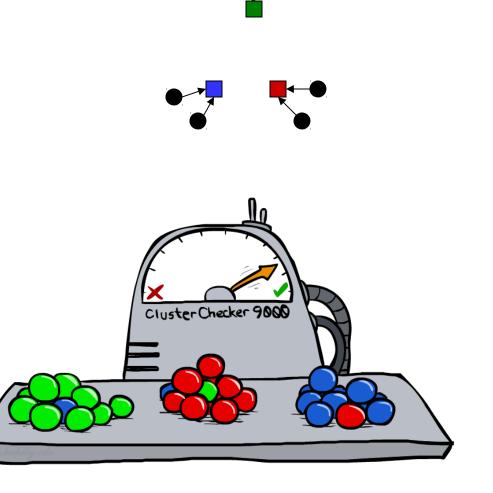


K-Means as Optimization

Consider the total distance to the means:

$$\phi(\{x_i\}, \{a_i\}, \{c_k\}) = \sum_i \operatorname{dist}(x_i, c_{a_i})$$
points from the means assignments

- Each iteration reduces phi
- Two stages each iteration:
 - Update assignments: fix means c, change assi
 - Update means: fix assignments a, change mea



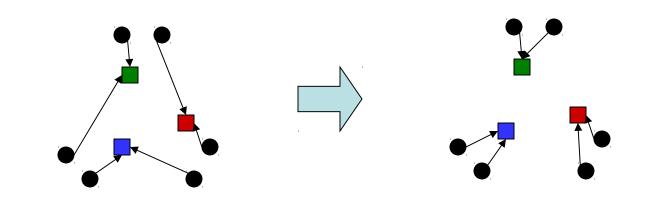
Phase I: Update Assignments

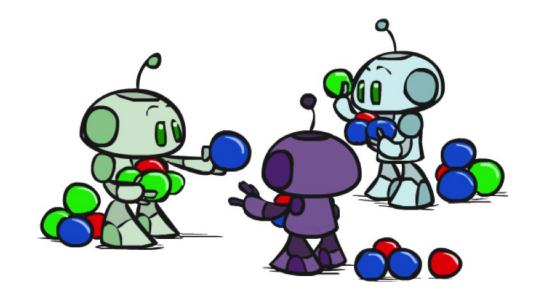
For each point, reassign to closest mean:

$$a_i = \underset{k}{\operatorname{argmin}} \operatorname{dist}(x_i, c_k)$$

Can only decrease total distance phi!

$$\phi(\{x_i\},\{a_i\},\{c_k\}) = \sum_i \operatorname{dist}(x_i,c_{a_i})$$



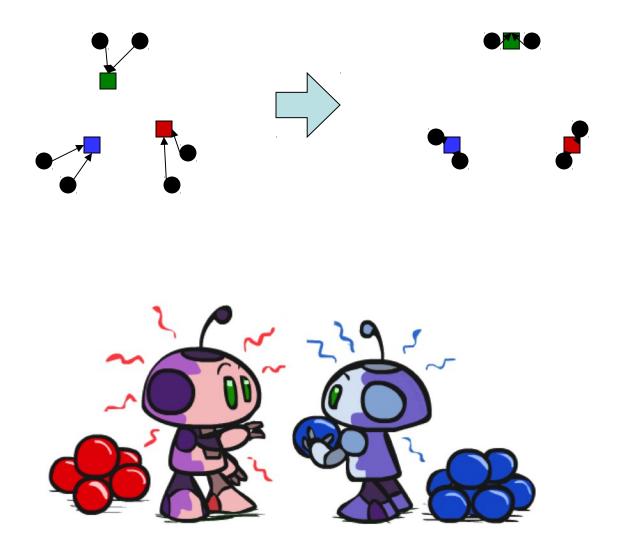


Phase II: Update Means

 Move each mean to the average of its assigned points:

$$c_k = \frac{1}{|\{i : a_i = k\}|} \sum_{i:a_i = k} x_i$$

- Also can only decrease total distance... (Why?)
- Fun fact: the point y with minimum squared Euclidean distance to a set of points {x} is their mean

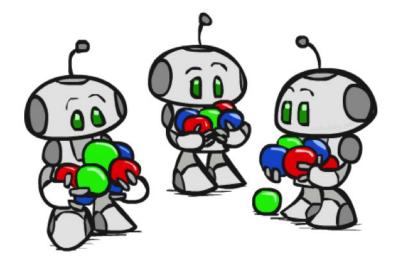


Initialization

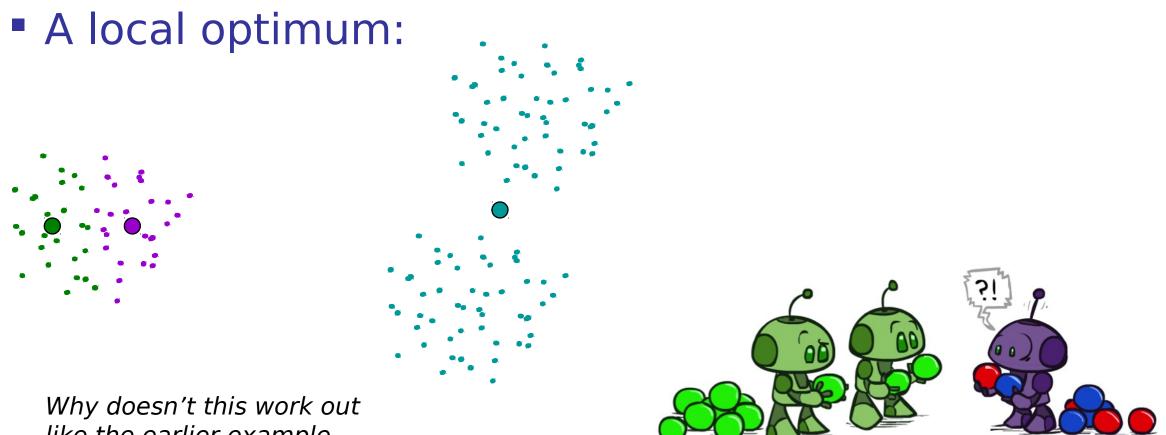
- K-means is non-deterministic
 - Requires initial means
 - It does matter what you pick!
 - What can go wrong?



 Various schemes for preventing this kind of thing: variancebased split / merge, initialization heuristics



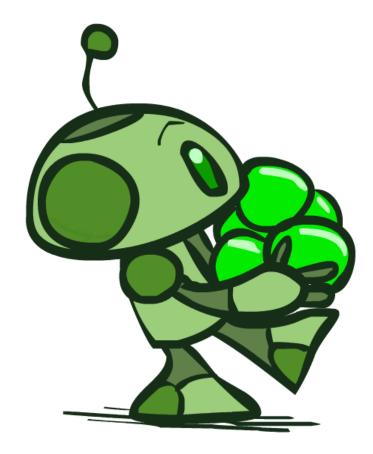
K-Means Getting Stuck



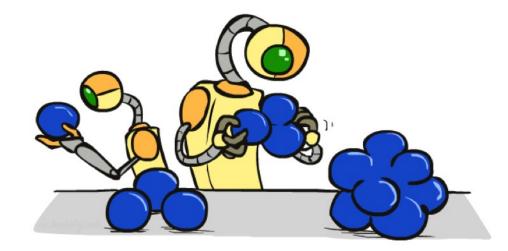
like the earlier example, with the purple taking over half the blue?

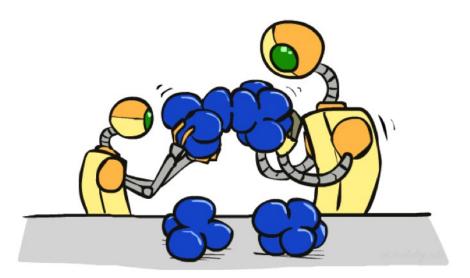
K-Means Questions

- Will K-means converge?
 - To a global optimum?
- Will it always find the true patterns in the data?
 - If the patterns are very very clear?
- Will it find something interesting?
- Do people ever use it?
- How many clusters to pick?



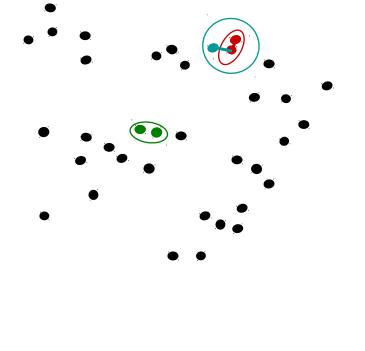
Agglomerative Clustering

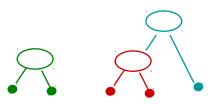




Agglomerative Clustering

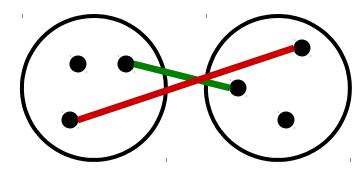
- Agglomerative clustering:
 - First merge very similar instances
 - Incrementally build larger clusters out of smaller clusters
- Algorithm:
 - Maintain a set of clusters
 - Initially, each instance in its own cluster
 - Repeat:
 - Pick the two closest clusters
 - Merge them into a new cluster
 - Stop when there's only one cluster left
- Produces not one clustering, but a family of clusterings represented by a dendrogram

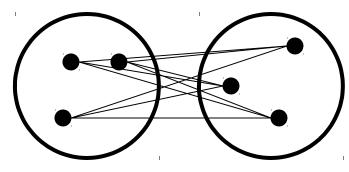


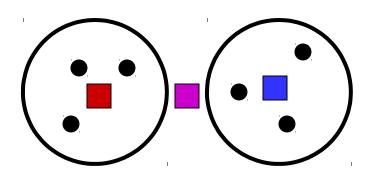


Agglomerative Clustering

- How should we define "closest" for clusters with multiple elements?
- Many options
 - Closest pair (single-link clustering)
 - Farthest pair (complete-link clustering)
 - Average of all pairs
 - Ward's method (min variance, like k-means)
- Different choices create different clustering behaviors







Example: Google News

