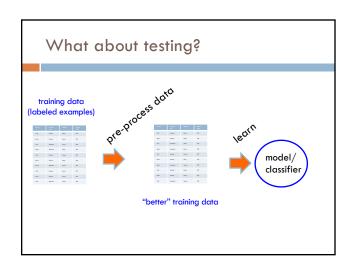
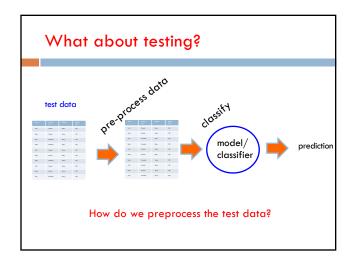


# So far... 1. Throw out outlier examples 2. Remove noisy features 3. Pick "good" features 4. Normalize feature values 1. center data 2. scale data (either variance or absolute) 5. Normalize example length 6. Finally, train your model!

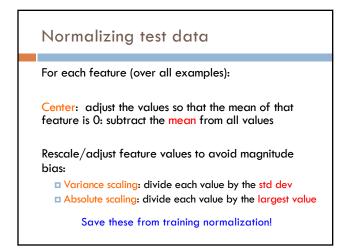


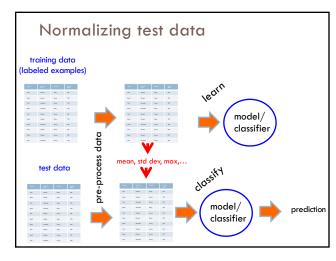


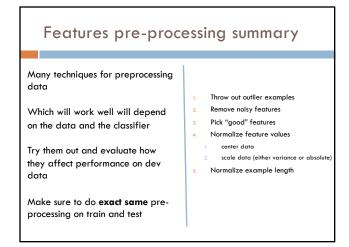
### 1. Throw out outlier examples 2. Remove noisy features 3. Pick "good" features 4. Normalize feature values 1. center data 2. scale data (either variance or absolute) 5. Normalize example length Which of these do we need to do on test data? Any issues?

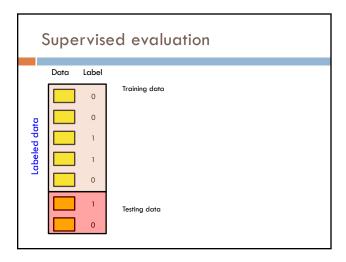
## Test data preprocessing Throw out outlier examples Remove irrelevant/noisy features Pick "good" features Normalize feature values Do these center data scale data (either variance or absolute) Normalize example length Whatever you do on training, you have to do the EXACT same on testing!

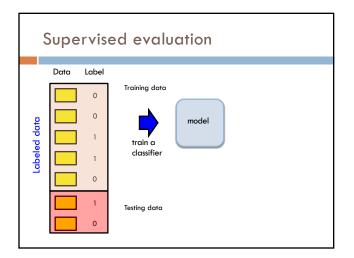
### Normalizing test data For each feature (over all examples): Center: adjust the values so that the mean of that feature is 0: subtract the mean from all values Rescale/adjust feature values to avoid magnitude bias: Variance scaling: divide each value by the std dev Absolute scaling: divide each value by the largest value What values do we use when normalizing testing data?

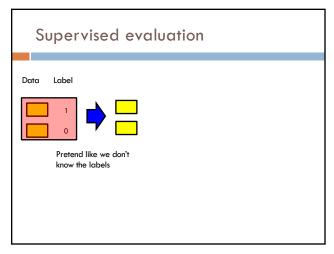


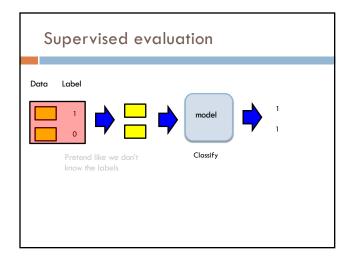


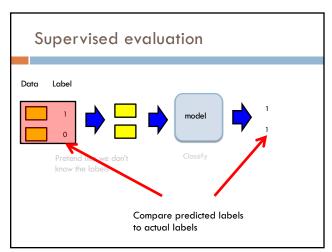


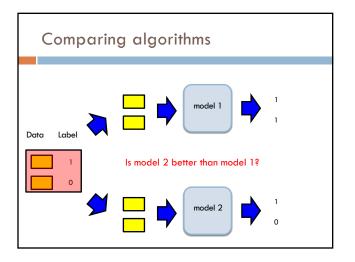


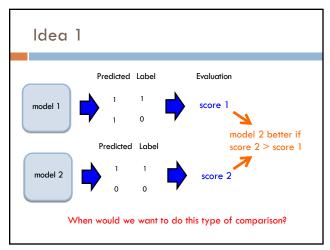


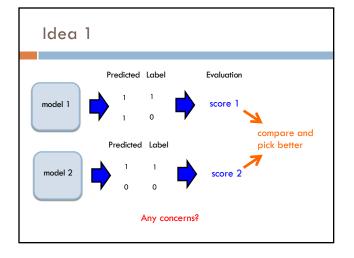


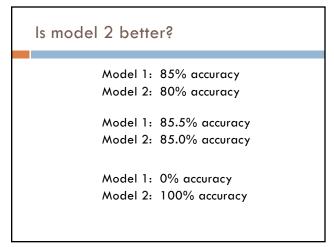










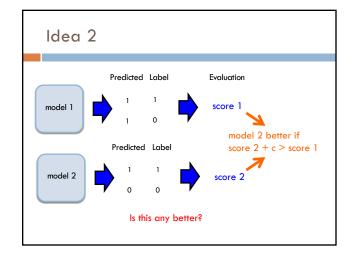


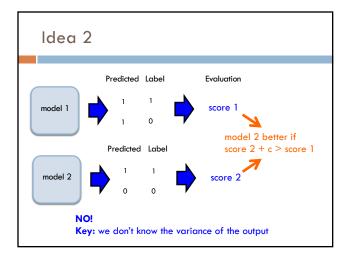
### Comparing scores: significance

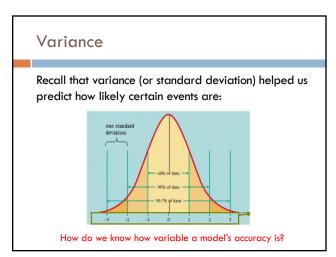
Just comparing scores on one data set isn't enough!

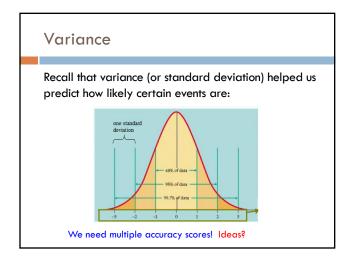
We don't just want to know which system is better on this particular data, we want to know if model 1 is better than model 2 in general

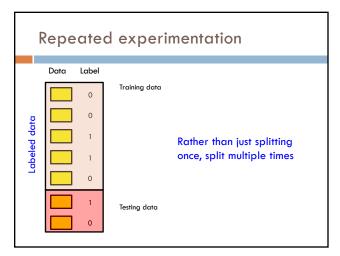
Put another way, we want to be confident that the difference is real and not just due to random chance

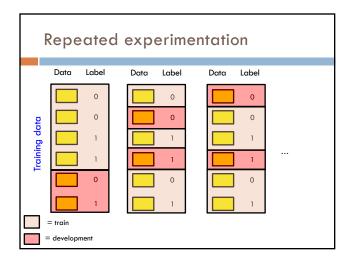


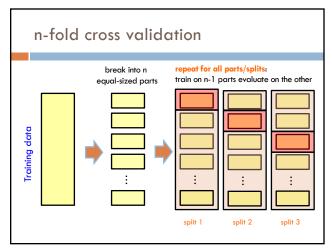


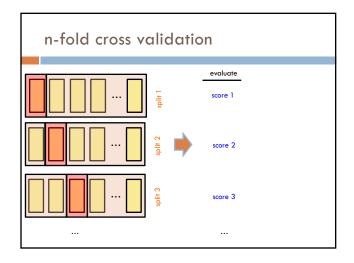












### n-fold cross validation

better utilization of labeled data

more robust: don't just rely on one test/development set to evaluate the approach (or for optimizing parameters)

multiplies the computational overhead by n (have to train n models instead of just one)

10 is the most common choice of n

### Leave-one-out cross validation

n-fold cross validation where n = number of examples

aka "jackknifing"

pros/cons?

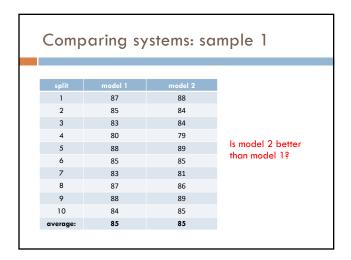
when would we use this?

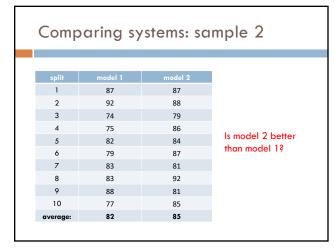
### Leave-one-out cross validation

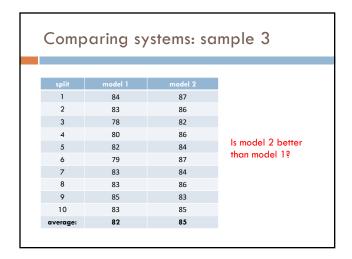
Can be very expensive if training is slow and/or if there are a large number of examples

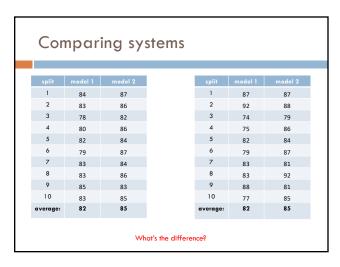
Useful in domains with limited training data: maximizes the data we can use for training

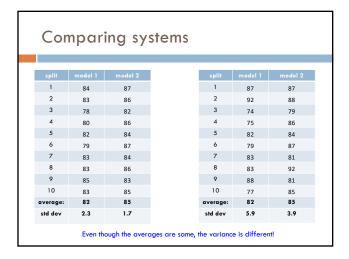
Some classifiers are very amenable to this approach (e.g.?)

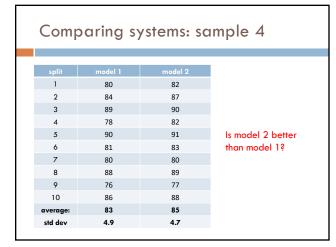


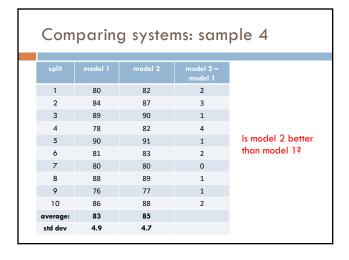


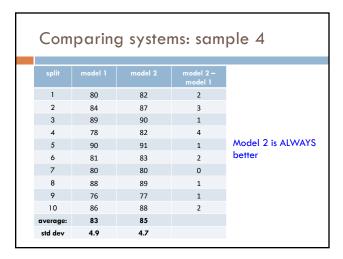












### Comparing systems: sample 4 80 82 84 87 89 90 82 How do we decide if 90 91 model 2 is better than model 1? 80 80 0 89 88 9 76 77 10 88 86 83 85 4.7 4.9 std dev

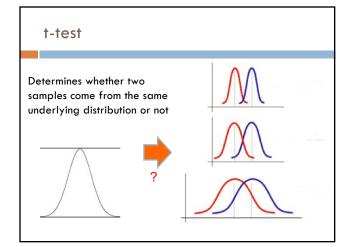
### Statistical tests

### Setup:

- Assume some default hypothesis about the data that you'd like to disprove, called the null hypothesis
- e.g. model 1 and model 2 are not statistically different in performance

### Test:

- Calculate a test statistic from the data (often assuming something about the data)
- Based on this statistic, with some probability we can reject the null hypothesis, that is, show that it does not hold



### t-test

Null hypothesis: model 1 and model 2 accuracies are no different, i.e. come from **the same** distribution

Assumptions: there are a number that often aren't completely true, but we're often not too far off

Result: probability that the difference in accuracies is due to random chance (low values are better)

### Calculating t-test

For our setup, we'll do what's called a "pair t-test"

- $\hfill\Box$  The values can be thought of as pairs, where they were calculated under the same conditions
- □ In our case, the same train/test split
- ☐ Gives more power than the unpaired t-test (we have more information)

For almost all experiments, we'll do a "two-tailed" version of the t-test

Can calculate by hand or in code, but why reinvent the wheel: use excel or a statistical package  $\,$ 

http://en.wikipedia.org/wiki/Student's\_t-test

### p-value

The result of a statistical test is often a p-value

p-value: the probability that the null hypothesis holds. Specifically, if we re-ran this experiment multiple times (say on different data) what is the probability that we would reject the null hypothesis incorrectly (i.e. the probability we'd be wrong)

Common values to consider "significant": 0.05 (95% confident), 0.01 (99% confident) and 0.001 (99.9% confident)

### Comparing systems: sample 1 2 85 84 84 80 79 Is model 2 better 5 88 89 than model 1? 85 85 83 81 They are the same with: 8 87 86

89

85

85

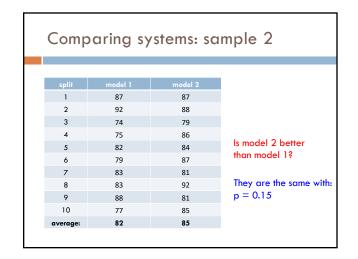
9

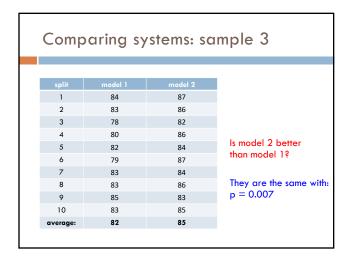
10

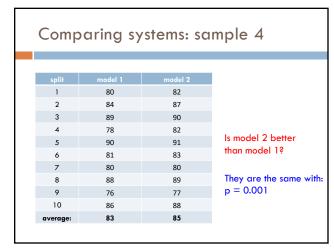
average:

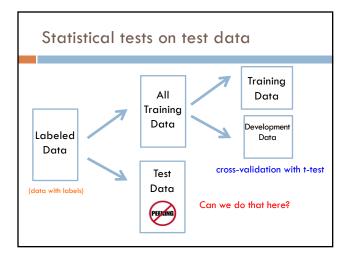
88

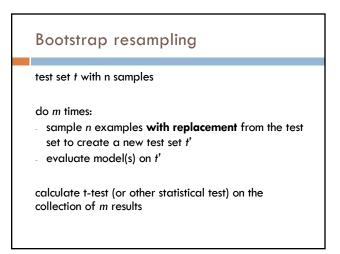
85

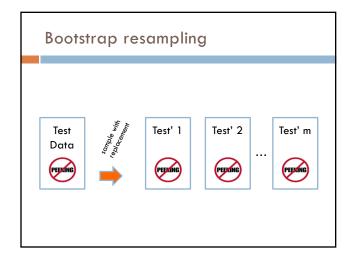


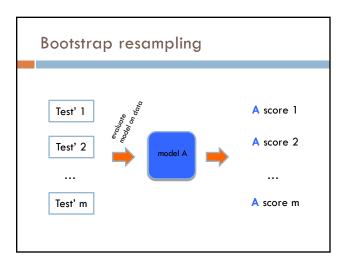


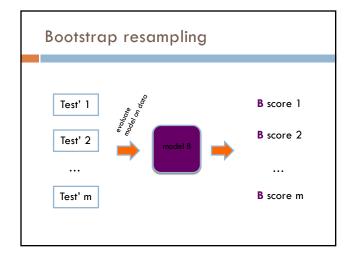


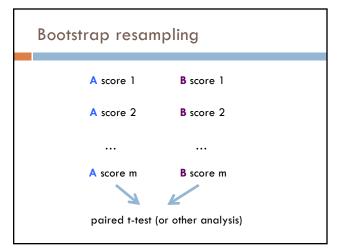












### Experimentation good practices

Never look at your test data!

### During development

- Compare different models/hyperparameters on development data
- use cross-validation to get more consistent results
- □ If you want to be confident with results, use a t-test and look for p = 0.05 (or even better)

For final evaluation, use bootstrap resampling combined with a t-test to compare final approaches