COLUMBIA MAILMAN SCHOOL UNIVERSITY of PUBLIC HEALTH



CROSS VALIDATION

Jeff Goldsmith, PhD Department of Biostatistics

Model selection

- When you have lots of possible variables, you have you choose which ones will go in your model
- In the best case, you have a clear hypothesis you want to test in the context of known confounders
- (Always keep in mind that no model is "true")

Model selection is hard

- Lots of times you're not in the best case, but still have to do something
- This isn't an easy thing to do
- For nested models, you have tests
 - You have to be worried about multiple comparisons and "fishing"
- For non-nested models, you don't have tests
 - AIC / BIC / etc are traditional tools
 - Balance goodness of fit with "complexity"

Questioning fit

- These are basically the same question:
 - Is my model not complex enough? Too complex?
 - Am I underfitting? Overfitting?
 - Do I have high bias? High variance?
- Another way to think of this is out-of-sample goodness of fit:
 Will my model generalize to future datasets?

COLUMBIA MAILMAN SCHOOL UNIVERSITY of PUBLIC HEALTH

Flexibility vs fit



Prediction accuracy

- Ideally, you could
 - Build your model given a dataset
 - Go out and get new data
 - Confirm that your model "works" for the new data
- That doesn't really happen
- So maybe just act like it does?

Cross validation

- Randomly split your data into "training" and "testing"
 - "Training" is data you use to build your model
 - "Testing" is data you use to evaluate out-of-sample fit
 - Exact ratio depends on data size, but I like 80 / 20
- Evaluate using root mean squared error :

$$RMSE = \sqrt{\frac{\sum_{i} (\hat{y}_{i} - y_{i})^{2}}{n}}$$



Refinements and variations

- Individual training / testing splits are subject to randomness
- Repeating the process
 - Illustrates variability in prediction accuracy
 - Can indicate whether differences in models are consistent across splits
- I usually repeat the training / testing split
- Folding (5-fold, 10-fold, k-fold, LOOCV) partitions data into equally-sized subsets
 - One fold is used as testing, with remaining folds as training
 - Repeated for each fold as testing
- I don't do this as often

Cross validation is general

- Can use to compare candidate models that are all "traditional"
- Comes up a lot in "modern" methods
 - Automated variable selection (e.g. lasso)
 - Additive models
 - Regression trees

Prediction as a goal

- In the best case, you have a clear hypothesis you want to test in the context of known confounders
 - I know I already said this, but it's important
- Prediction accuracy matters as well
 - Different goal than statistical significance
 - Models that make poor predictions probably don't adequately describe the data generating mechanism, and that's bad

Tools for CV

- Lots of helpful functions in modelr
 - add_predictions() and add_residuals()
 - rmse()
 - crossv_mc()
- Since repeating the process can help, list columns and map come in handy a lot too :-)

