Model selection

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Model selection problems

Model selection problem: how to select the f() that maps features X to target y? We'll look at two examples of model selection problems, but there are many more.

Choosing model complexity

We need to select a model of appropriate complexity -

- · what does that mean, and
- · how do we select one?

Model order selection problem

- Given data $(x_i, y_i), i = 1 \cdots, N$ (one feature)
- Polynomial model: $\hat{y} = w_0 + w_1 x + \dots + w_d x^d$
- d is degree of polynomial, called model order
- Given d, can get regression coefficients via OLS
- Model order selection problem: choosing d

Using loss function for model order selection?

Suppose we would "search" over each possible d:

- Fit model of order d on training data, get ${\bf w}$
- Compute predictions on training data: $\hat{y}_i, i = 1, ..., n$.
- Compute loss function (e.g. RSS) on training data: $RSS = \sum_{i=1}^n (y_i \hat{y_i})^2$
- ullet Select d that minimizes loss
- Problem: loss function always decreasing with d (training error decreases with model complexity!)

Feature selection problem

- Linear model: $\hat{y} = w_0 + w_1 x_1 + \dots + w_d x_d$
- Model target y as a function of features $\mathbf{x} = (x_1, \cdots, x_d)$
- · Many features, only some are relevant
- Feature selection problem: fit a model with a small number of features

Why use a subset of features?

- High risk of overfitting if you use all features!
- For linear regression, there's a unique OLS solution only if n > d
- For linear regression, when $N \geq p$, variance increases linearly with number of parameters, inversely with number of samples. (Not derived in class, but read extra notes posted after class at home.)
- Important applications where you have many features:
 - EEG measure brain activity with electrodes, typically >10,000 "voxels" but only 100s of observations
 - DNA microarray data measures "expression" levels of large number of genes (~1000) but only a small number of data points (~100)

Feature selection problem - formal

Problem: given high dimensional data $\mathbf{X} \in R^{n \times d}$ and target variable y, Select a subset of k << d features, $\mathbf{X}_S \in R^{n \times k}$ that is most relevant to target y.

Cross validation

Simple train/validation/test split

- Divide data into training, validation, test sets
- · For each candidate model, learn model parameters on training set
- Measure error for all models on validation set
- · Select model that minimizes error on validation set
- · Evaluate model on test set

Note: sometimes you'll hear "validation set" and "test set" used according to the reverse meanings.

Simple train/validation/test algorithm

- Get data X, y and split into training, validation, and test.
- Loop over models of increasing complexity: For $p=1,\ldots,p_{max}$
 - Fit: $\hat{w}_p = \mathrm{fit}_p(X_{tr}, y_{tr})$
 - Predict: $\hat{y}_{v,p} = \operatorname{pred}(X_v, \hat{w}_p)$
- Score: $S_p = \text{score}(y_v, \hat{y}_{v,p})$ Select model order with best score: $p^* = \operatorname{argmin}_p S_p$
- Evaluate: $S_{p^*} = \text{score}(y_{ts}, \hat{y}_{ts,p^*})$ where $\hat{y}_{ts,p^*} = \text{pred}(X_{ts}, \hat{w}_{p^*})$

Problems with simple split

- Fitted model (and test error!) varies a lot depending on samples selected for training and validation.
- Fewer samples available for estimating parameters.
- Especially bad for problems with small number of samples.

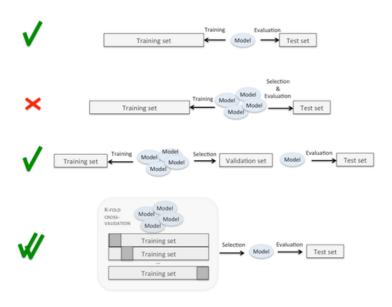


Figure 1: Summary of approaches. Source.

K-fold cross validation

Alternative to simple split:

- Divide data into K equal-sized parts (typically 5, 10)
- For each of the "splits": evaluate model using K-1 parts for training, last part for validation
- ullet Average the K validation scores and choose based on average

K-fold CV illustrated

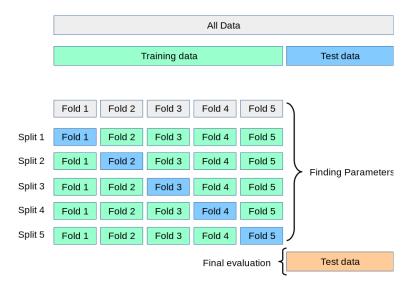


Figure 2: K-fold CV

K-fold CV - pseudocode (1)

Outer loop over folds: for i=1 to K

- Split training data into training and validation:
- Inner loop over models of increasing complexity: For p=1 to p_{max} ,

 - $$\begin{split} & \textbf{- Fit: } \hat{w}_{p,i} = \text{fit}_p(X_{tr_i}, y_{tr_i}) \\ & \textbf{- Predict: } \hat{y}_{v_i,p} = \text{pred}(X_{v_i}, \hat{w}_{p,i}) \\ & \textbf{- Score: } S_{p,i} = score(y_{v_i}, \hat{y}_{v_i,p}) \end{split}$$

K-fold CV - pseudocode (2)

- Find average score (across K scores) for each model: $ar{S}_p$
- Select model with best average score: $p^* = \operatorname{argmin}_p \bar{S}_p$
- Re-train model on entire training set: $\hat{w}_{p^*} = \text{fit}_p(\hat{X_{tr}}, y_{tr})$
- · Evaluate new fitted model on test set

K-fold CV - how to split?

Choose the split strategy based on the data:

· Shuffle and split ("standard" K-fold CV)

- Split without shuffling
- · Stratified K-fold CV: make sure distribution of target variable is similar in each part
- Group K-fold CV: samples from the same "group" go in either training or validation data, but never in both
- · Time series CV

Selecting the right K-fold CV is very important for avoiding data leakage!

Refer to the function documentation for more examples.

One standard error rule

- Model selection that minimizes mean error often results in too-complex model
- · One standard error rule: use simplest model where mean error is within one SE of the minimum mean error

One standard error rule - algorithm (1)

- Given data X,y Compute score $S_{p,i}$ for model p on fold i (of K)
- Compute average (\bar{S}_p) , standard deviation σ_p , and standard error of scores:

$$SE_p = \frac{\sigma_p}{\sqrt{K-1}}$$

One standard error rule - algorithm (2)

"Best score" (smallest loss) model selection:

$$p^* = \operatorname*{argmin}_p \bar{S}_p$$

One SE rule:

Compute target score: $S_t = \bar{S}_{p^*} + SE_{p^*}$

then select simplest model with score lower than target:

$$p^{*,\mathrm{1SE}} = \min\{p | \bar{S}_p \leq S_t\}$$

Note: this assumes you are using a "smaller is better" metric such as MSE. If you are using a "larger is better" metric, like R2, adjust the algorithm accordingly.

TODO: add illustration

TODO: add explanation of how to adjust to a "larger is better" metric.