

# Criteria for model comparison

- In the last video, during each step of the forward, backward, and stepwise selection, we are using an  $F$ -test to choose between two models
- We can use other criteria to compare between models
- There are many proposed approach in the literature
- They have their own advantages and limitations
- But there is no 'standard' or 'best' criterion to use
- We'll look at some popular model selection criteria

# Criteria for comparing models

There are a number of commonly used criteria for comparing models:

- $R^2$
- Adjusted  $R^2$
- Mallow's  $C_p$
- Akaike information criterion (AIC)
- Bayesian information criterion (BIC)
- PRESS statistics

# $R^2$

The proportion of variation in  $Y$  (about  $\hat{Y}$ ) explained by the predictors in the model is known as the coefficient of multiple determination, denoted by  $R^2$ ,

$$R^2 = \frac{SSR}{SST} = 1 - \frac{SSE}{SST}.$$

*Handwritten red annotations:*  
-  $SSR$  is labeled  $SSE_D$  with a red line pointing to it.  
-  $SSE$  is labeled  $SSE_F$  with a red line pointing to it.  
-  $SST$  is labeled  $SSE_R$  with a red line pointing to it.

- larger  $R^2$  is preferred
- Useful for comparing models with same number of predictors

*In this case, we are effectively choosing the model with the smallest SST*

# Adjusted $R^2$

The adjusted  $R^2$  is the  $R^2$  'corrected' for degrees of freedom:

$$\tilde{R}^2 = 1 - \frac{MSE}{MST} = 1 - \left( \frac{n-1}{n-p} \right) (1 - R^2).$$

*Handwritten notes:  $MSE_F$  above the fraction,  $MSE_R$  below the fraction.*

- larger  $\tilde{R}^2$  is preferred
- can compare models of different size

$$\tilde{R}^2 = 1 - \frac{MSE}{MST} = 1 - \frac{\frac{SSE}{n-p}}{\frac{SST}{n-1}} = 1 - \left( \frac{n-1}{n-p} \right) \left( \frac{SSE}{SST} \right) = 1 - \left( \frac{n-1}{n-p} \right) (1 - R^2)$$

# Mallow's $C_p$

The Mallow's  $C_p$  statistic compares the predictive ability of subset models to that of the full model:

$$C_p = \frac{SSE}{\hat{\sigma}^2} - (n - 2p),$$

*smaller model* (pointing to SSE)  
*MSE of full model =  $\sigma^2$*  (pointing to  $\hat{\sigma}^2$ )

where  $\hat{\sigma}^2$  is an estimate of  $\sigma^2$  based on the full model.

- the model with the smallest  $p$  for which  $C_p \approx p$

$E(SSE) = (n-p)\sigma^2$  if the smaller model is adequate

$E(SSE) > (n-p)\sigma^2$  if the smaller model is not adequate

$$E[C_p] = E\left(\frac{SSE}{\hat{\sigma}^2}\right) - (n-2p) \approx \frac{(n-p)\sigma^2}{\sigma^2} - (n-2p) = (n-p) - (n-2p) = p$$

# Akaike information criterion (AIC)

The Akaike information criterion (AIC) is defined as

$$AIC = 2p - 2\ln(\hat{L}),$$

where  $p$  is the number of parameter in the model, and  $\ln(\hat{L})$  is the log likelihood evaluated at the maximum likelihood estimates.

- the model with the lowest AIC is preferred

For linear regression,  $\ln(\hat{L}) = \text{constant} - \frac{n}{2} \ln(\text{SSE})$

$$AIC = \text{constant} + 2p + n \ln(\text{SSE})$$

# Akaike information criterion corrected (AICc)

To adjust for small sample sizes, the AICc is used. It is defined as

$$AICc = AIC + \overset{\text{penalty term}}{\boxed{\frac{2p(p+1)}{n-p-1}}},$$

where  $n$  is the sample size.

As  $n \rightarrow \infty$ , penalty term  $\rightarrow 0$ ,  $AICc \rightarrow AIC$

# Bayesian information criterion (BIC)

A more stringent criterion with respect to the number of parameters is the Bayesian information criterion (BIC). It is defined as

$$BIC = \boxed{\ln(n)}p - 2 \ln(\hat{L}).$$

- the model with the lowest BIC is preferred

For linear regression,  $BIC = p \ln(n) + n \ln(\hat{L})$

In R, the `step()` function uses the formula

$$IC = \boxed{k}p - 2 \ln(\hat{L})$$

It uses AIC by default.

For AIC, set  $k=2$ .

For BIC, set  $k = \ln(n)$ .

If we want to use P-value with an arbitrary significance level  $\alpha$ , set  $k = \chi^2_{1,\alpha}$ .



## Example 4.4

Consider again the marks data in Example 4.1.

Fit a multiple linear regression to the data using AIC and

- (a) Forward selection
- (b) Backward selection
- (c) Stepwise selection

# Example 4.4 Solutions

If using BIC:  $k = \log(n)$

```
fAIC <- step(null, scope=scope, direction="forward")
```

```
## Start: AIC=-938.43
```

```
## E ~ 1
```

```
##
```

##		Df	Sum of Sq	RSS	AIC
##	+ A6	1	9.3016	11.853	-1132.80
##	+ A4	1	7.1621	13.993	-1076.55
##	+ OQ	1	7.1258	14.029	-1075.67
##	+ A5	1	6.9001	14.255	-1070.26
##	+ A3	1	4.2472	16.908	-1012.40
##	+ A2	1	1.7407	19.414	-965.54
##	+ A1	1	1.0852	20.070	-954.28
##	<none>			21.155	-938.43

```
##
```

```
## Step: AIC=-1132.8
```

```
## E ~ A6
```

```
##
```

##		Df	Sum of Sq	RSS	AIC
##	+ OQ	1	1.12811	10.725	-1164.7
##	+ A3	1	0.56043	11.293	-1147.2
##	+ A4	1	0.55393	11.299	-1147.0
##	+ A5	1	0.32088	11.532	-1140.1
##	+ A2	1	0.09538	11.758	-1133.5
##	+ A1	1	0.08001	11.773	-1133.1
##	<none>			11.853	-1132.8

backward selection: "backward"

stepwise selection: "both"

add A6

Find the lowest AIC value.

Add the corresponding variable to our model.

add OQ

# Example 4.4 Solutions (cont.)

```
##
## Step: AIC=-1164.71
## E ~ A6 + OQ
##
```

	Df	Sum of Sq	RSS	AIC
## + A3	1	0.33372	10.391	-1173.4
## + A4	1	0.18839	10.537	-1168.7
## + A5	1	0.09645	10.629	-1165.8
## <none>			10.725	-1164.7
## + A1	1	0.03104	10.694	-1163.7
## + A2	1	0.02419	10.701	-1163.5

add A3

```
##
## Step: AIC=-1173.43
## E ~ A6 + OQ + A3
##
```

	Df	Sum of Sq	RSS	AIC
## + A4	1	0.070882	10.320	-1173.8
## <none>			10.391	-1173.4
## + A5	1	0.039884	10.351	-1172.7
## + A2	1	0.006228	10.385	-1171.6
## + A1	1	0.000108	10.391	-1171.4

add A4

# Example 4.4 Solutions (cont.)

```
##  
## Step: AIC=-1173.75  
## E ~ A6 + OQ + A3 + A4  
##  
##      Df Sum of Sq  RSS   AIC  
## <none>      10.320 -1173.8  
## + A5       1 0.0235297 10.297 -1172.5  
## + A2       1 0.0136998 10.307 -1172.2  
## + A1       1 0.0003186 10.320 -1171.8
```

Handwritten annotations: A red box highlights the model formula and the first row of the table. A green box highlights the rest of the table. A green arrow points from the text "current model" to the "<none>" entry in the first row.

The current model <none> has the lowest AIC. Hence we can stop here.

# Example 4.4 Solutions (cont.)

```
summary(fAIC)
```

```
##
## Call:
## lm(formula = E ~ A6 + OQ + A3 + A4, data = stats_marks)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.81971 -0.06460  0.02923  0.09030  0.61607
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   0.12306    0.03321   3.705 0.000247 ***
## A6            0.32917    0.04666   7.054 1.00e-11 ***
## OQ            0.18475    0.03868   4.777 2.67e-06 ***
## A3            0.12196    0.04611   2.645 0.008549 **
## A4            0.08014    0.05291   1.515 0.130826
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.1758 on 334 degrees of freedom
## Multiple R-squared:  0.5121, Adjusted R-squared:  0.5063
## F-statistic: 87.66 on 4 and 334 DF,  p-value: < 2.2e-16
```

# PRESS statistic

The prediction sum of squares (PRESS) is used to assess a model's predictive ability, and is given by

$$PRESS = \sum_{i=1}^n (y_i - \hat{y}_{(i)})^2,$$

where  $\hat{y}_{(i)}$  is the predicted value of  $y_i$  using the model fitted with  $i$ th observation removed.

- smaller PRESS is preferred

## Model validation

Q: Is our model generalizable beyond our sample data?

Ideally, we want to get new observations to validate our model:

(1) Refit the model with the new observations included.

Check to see if the parameter of the new model is significantly different from our previous model.

(2) Use our original model to predict the response of the new observations. Calculate the prediction errors for the new observations.

In practice, it may be difficult to get new observations. An alternative is to split our data into a training set and a validation/testing set. This is called cross-validation (CV).

(1) Fit our model using the training set

(2) Calculate prediction error using the testing set.

# Cross-validation

Another useful method to assess how good a model is for prediction is cross-validation.

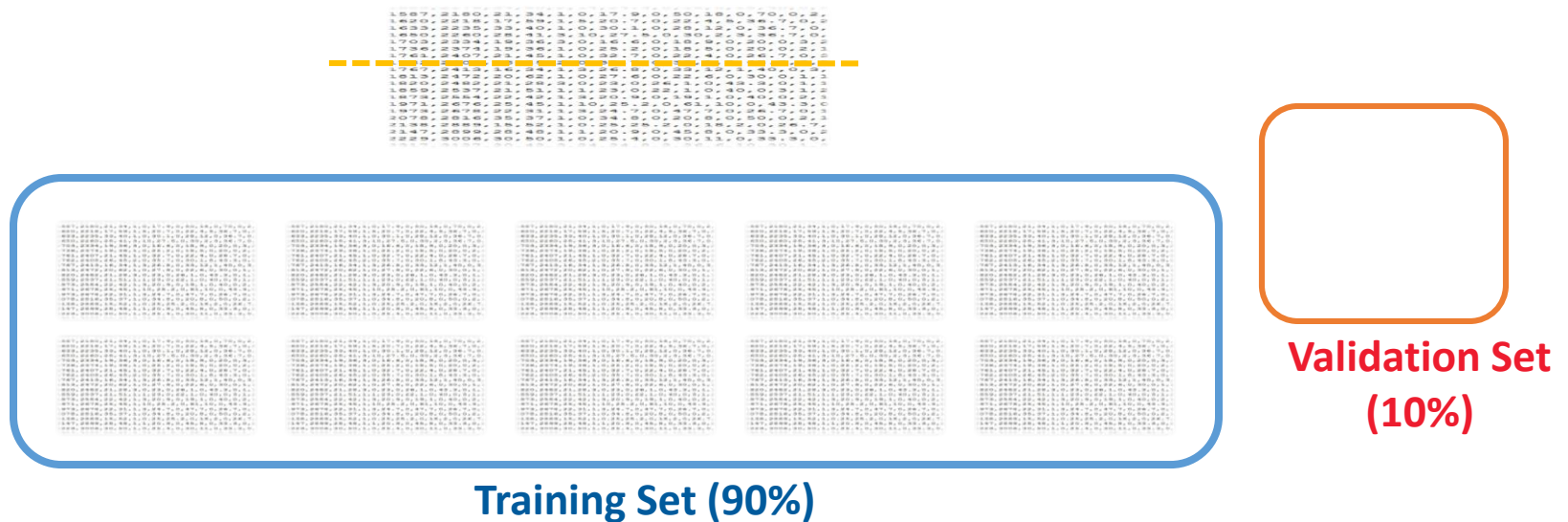
For  $k$ -fold cross-validation, you split the data into  $k$  parts.

In each step, train the model on  $k - 1$  parts and test for  $k$ th part.



# Cross-validation

- 10-fold cross-validation



- Training set: for model development
- Validation set: to assess the accuracy on new data
- Repeat 10 times: estimate prediction error

# Notation

Label each part

$$C_1, C_2, \dots, C_K$$

Let the number of observations in  $C_k$  be  $n_k$  ( $k = 1, 2, \dots, K$ ) so that

$$n = \sum_{k=1}^K n_k ,$$

i.e.  $n$  is the total number of observations.

# Prediction error

The cross validation estimate of the prediction error is

$$CV_{(k)} = \sum_{k=1}^K \frac{n_k}{n} MSE_k,$$

weighted sum  
of the MSE of  
the  $k$  folds

where

$$MSE_k = \sum_{i \in C_k} \frac{(y_i - \hat{y}_i)^2}{n_k},$$

where  $\hat{y}_i$  is the fitted value for observation  $i$  for the model with part  $k$  removed.

If  $k = n$ , we have leave-one-out cross validation (LOOCV). In this case,  $CV(n) = \text{PRESS}$ .