

Stat 206: Linear Models

Lecture 15

Nov. 20, 2019

Recap: Key Components for Model Selection

- **Criterion to compare models:**
 - R_a^2 , C_p , AIC_p , BIC_p , $Press_p$, etc.
- **Procedure to search for good model(s):**
 - *Best subset selection*: Exhaustive search; When the number of potential X variables is not too big
 - *Stepwise regression*: Greedy search; The number of potential X variables can be large.

Surgical Unit: Model Selection Criteria

Consider X_1, X_2, X_3, X_4 (clotting, prognostic, enzyme, liver) as the potential pool of X variables. There are 16 sub-models.

p	intercept	X1	X2	X3	X4	sse	R ²	R ² _a	Cp	aic	bic	press
1	1	0	0	0	0	12.805	0.000	0.000	151.569	-75.716	-73.727	13.292
2	1	0	0	1	0	7.334	0.427	0.416	66.518	-103.811	-99.833	8.329
2	1	0	0	0	1	7.408	0.421	0.410	67.696	-103.268	-99.290	8.024
2	1	0	1	0	0	9.974	0.221	0.206	108.469	-87.205	-83.227	10.738
2	1	1	0	0	0	12.028	0.061	0.043	141.093	-77.096	-73.118	13.508
3	1	0	1	1	0	4.313	0.663	0.650	20.523	-130.479	-124.512	5.066
3	1	0	0	1	1	5.132	0.599	0.583	33.536	-121.089	-115.122	6.123
3	1	1	0	1	0	5.783	0.548	0.531	43.873	-114.644	-108.677	6.989
3	1	0	1	0	1	6.620	0.483	0.463	57.175	-107.342	-101.375	7.474
3	1	1	0	0	1	7.299	0.430	0.408	67.961	-102.070	-96.103	8.472
3	1	1	1	0	0	9.437	0.263	0.234	101.937	-88.194	-82.227	11.055
4	1	1	1	1	0	3.109	0.757	0.743*	3.388*	-146.161*	-138.205*	3.914*
4	1	0	1	1	1	3.615	0.718	0.701	11.434	-138.011	-130.055	4.598
4	1	1	0	1	1	4.970	0.612	0.589	32.960	-120.823	-112.867	6.209
4	1	1	1	0	1	6.568	0.487	0.456	58.358	-105.763	-97.807	7.902
5	1	1	1	1	1	3.084	0.759*	0.739	5.000	-144.587	-134.642	4.069

Within each subset size, models are sorted in ascending SSE .

Consequently, within each subset size, $R_p^2, R_{a,p}^2$ are from the largest to the smallest and C_p, BIC_p, AIC_p are from the smallest to the largest. $Press_p$ may not be monotone with SSE .

AIC_p and BIC_p Criteria

- *Akaike's information criterion (AIC):*

$$AIC_p = n \log \frac{SSE_p}{n} + 2p.$$

- *Bayesian information criterion (BIC):*

$$BIC_p = n \log \frac{SSE_p}{n} + (\log n)p.$$

- **We should look for models with small AIC (BIC).**
 - Surgical unit. The model with X_1, X_2, X_3 has the smallest AIC and BIC among the models being considered.

- The first term: $n \log \frac{SSE_p}{n}$ reflects the of the model to the observed data.
 - It by adding more X variables into the model.
- The second term, $2p$ for AIC and $(\log n)p$ for BIC, reflects
 - It by adding more X variables into the model.
 - If $n \geq 8$, then $\log n > 2$ and BIC puts penalty on model complexity and tends to choose models than AIC.

- The first term: $n \log \frac{SSE_p}{n}$ reflects the *goodness-of-fit* of the model to the **observed data**.
 - It decreases by adding more X variables into the model.
- The second term, $2p$ for AIC and $(\log n)p$ for BIC, reflects model complexity.
 - It increases by adding more X variables into the model.
 - If $n \geq 8$, then $\log n > 2$ and BIC puts more penalty on model complexity and tends to choose smaller models than AIC.

- Overly simplified models have low model complexity (p), but they tend to have high SSE (underfitting; high bias).
- Overly complicated models may have a high SSE, but they have high model complexity (overfitting, high variance).
- By minimizing AIC (or BIC), we are trying to find a model that balances between model complexity and the goodness-of-fit.

- Overly simplified models have small model complexity (p), but they tend to have large SSE (underfitting; high bias).
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$Press_p$ Criterion

Predicted residual sum of squares ($Press_p$):

$$Press_p = \sum_{i=1}^n (Y_i - \hat{Y}_{i(i)})^2.$$

- Y_i is the observed response of the i th case.
- $\hat{Y}_{i(i)}$ is the predicted value for the i th case obtained by fitting the model only using $n - 1$ cases excluding case i .
- $Press_p$ is also known as *leave-one-out-cross-validation* (LOOCV).
- Models with small $Press_p$ are considered good in terms of predictive ability.
 - Surgical unit: the model with X_1, X_2, X_3 has $Press_p = 3.914$ which is the smallest among all models being considered here.

Calculate $Press_p$

$Press_p$ can be calculated without actually performing n regressions.

- This is because the *deleted residual* for the i th case:

$$d_i := Y_i - \widehat{Y}_{i(i)} = \quad , \quad i = 1, \dots, n.$$

where $e_i = Y_i - \widehat{Y}_i$ is the residual of the i th case and h_{ii} is the i th diagonal element of the hat matrix \mathbf{H} , both from the regression fit using .

- So

Calculate $Press_p$

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- This is because the *deleted residual* for the i th case:

$$d_i := Y_i - \widehat{Y}_{i(i)} = \frac{e_i}{1 - h_{ii}}, \quad i = 1, \dots, n.$$

where $e_i = Y_i - \widehat{Y}_i$ is the residual of the i th case and h_{ii} is the i th diagonal element of the hat matrix \mathbf{H} , both from the regression fit using **all** n cases.

- So

$$Press_p = \sum_{i=1}^n \frac{(Y_i - \widehat{Y}_i)^2}{(1 - h_{ii})^2}.$$

Derive the Deleted Residuals

Optional Reading.

- Define $\tilde{\mathbf{Y}}$ by replacing the i th element of the response vector \mathbf{Y} with the leave- i -out predicted value $\hat{Y}_{i(i)}$ of the i th case:

$$\tilde{\mathbf{Y}} = (Y_1, \dots, Y_{i-1}, \hat{Y}_{i(i)}, Y_{i+1}, \dots, Y_n)^T.$$

- Let $\hat{\beta}_{(i)}$ be the leave- i -out LS fitted regression coefficients. Then $\hat{\beta}_{(i)}$ is also the LS fitted regression coefficients by using $\tilde{\mathbf{Y}}$ as the response vector, i.e. $\hat{\beta}_{(i)} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \tilde{\mathbf{Y}}$. *Why?*
- The leave- i -out fitted values are:

$$\hat{\mathbf{Y}}_{(i)} = \mathbf{X} \hat{\beta}_{(i)} = H \tilde{\mathbf{Y}} = H(\mathbf{d}_{(i)} + \mathbf{Y}), \quad \mathbf{d}_{(i)} = \tilde{\mathbf{Y}} - \mathbf{Y} = (0, \dots, -d_i, \dots, 0)^T.$$

- Subtracting the i th element from Y_i on both sides gives:

$$d_i = h_{ii} d_i + e_i \implies d_i = \frac{e_i}{1 - h_{ii}}.$$

Surgical Unit: Full Model X_1, X_2, X_3, X_4

```
> fit.f = lm(log(Y) ~ X1 + X2 + X3 + X4, data = data.o)
> summary(fit.f)
Call:
lm(formula = log(Y) ~ X1 + X2 + X3 + X4, data = data.o)
...
Coefficients:
Estimate Std. Error t value Pr(>|t|)
(Intercept) 3.851933    0.266263   14.467 < 2e-16 ***
X1           0.083739    0.028834    2.904 0.00551 **
X2           0.012671    0.002315    5.474 1.50e-06 ***
X3           0.015627    0.002100    7.440 1.38e-09 ***
X4           0.032056    0.051466    0.623 0.53627
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.2509 on 49 degrees of freedom
Multiple R-squared: 0.7591,    Adjusted R-squared: 0.7395
F-statistic: 38.61 on 4 and 49 DF,  p-value: 1.398e-14
> anova(fit.f)
Analysis of Variance Table
```

```
Response: log(Y)
Df Sum Sq Mean Sq F value    Pr(>F)
X1      1  0.7770   0.7770  12.3443 0.0009618 ***
X2      1  2.5904   2.5904  41.1565 5.341e-08 ***
X3      1  6.3286   6.3286 100.5490 1.838e-13 ***
X4      1  0.0244   0.0244   0.3879 0.5362698
Residuals 49 3.0841   0.0629
```

Surgical Unit: Full Model

- Full model has $P = 5$ and

$$SSE = 3.0841, \text{ MSE} = 0.0629, R^2 = 0.7591, R_a^2 = 0.7395.$$

- By definition, for the full model, $C_P = P = 5$.
- Sample size $n = 54$, so for the full model:
 $AIC_P = 54 \log(3.0841/54) + 2 \times 5 = -144.5871$ and
 $BIC_P = 54 \log(3.0841/54) + \log(54) \times 5 = -134.6422$.
- $Press_p = 4.069$.

```
> e.f=fit.f$residuals  ## residuals  
> h.f=influence(fit.f)$hat  ## diagonals of hat matrix  
> press.f= sum(e.f^2/(1-h.f)^2)  ## calculate press
```

Model Search Procedures

- The number of possible models, 2^{P-1} , grows very fast with the number potential X variables $P - 1$.
- Evaluating every possible model can be computationally infeasible even for moderate P .
- A variety of search procedures have been developed to efficiently search for the “best” model(s) in the model space.
 - **Stepwise regression procedures**
 - Best subsets algorithms: Not applicable when the pool of potential X variables is large.

Stepwise Regression Procedures

- Applicable to situations with a large number of potential X variables.
- Use “greedy” search strategies by developing a sequence of models, at each step adding or deleting only one X variable according to a pre-specified criterion (e.g., AIC).
- May end up with a *suboptimal model* rather than the global “best” model.
- Commonly used stepwise procedures include: *forward stepwise*, *forward selection*, *backward stepwise* and *backward elimination*.

Forward Stepwise Procedure

Need to specify:

- A model selection criterion, e.g., *AIC*.
- An initial model M_0 , usually a small model, e.g., the null-model with no X variable.
- The pool of potential X variables \mathcal{X} .
- The set of X variables that will always be in the model \mathcal{X}_0 , e.g., the intercept term.

Starting from the initial model M_0 , at each step:

- (a) Consider the X variables in the potential pool X that are not currently in the model. Examine the change in the criterion by adding each such variable into the current model.
- (b) Consider the X variables that are already in the model but not in the set X_0 . Examine the change in the criterion by dropping each such variable out of the current model.
- Choose the operation that improves the criterion the most and update the current model accordingly. Repeat steps (a) and (b) for the updated model.
- If there is no operation that can improve the criterion anymore, then stop the search procedure and return the current model as the selected model.

Forward Selection and Backward Elimination

- Forward selection is a simplified version of forward stepwise procedure, omitting the considerations of dropping a variable currently in the model at each step.
- Backward elimination is the opposite of the forward selection.
 - It starts with a “big” initial model, e.g., the full model.
 - At each step, it examines the change of the criterion by dropping a variable currently in the model.
- Backward stepwise procedure. *Guess what is it?*
- Another commonly used strategy is to perform one pass of forward selection followed by one pass of backward elimination.

stepAIC () Function

We can use the `stepAIC()` function in the `MASS` library to perform various stepwise regression.

- `direction='both'` corresponds to forward stepwise procedure or backward stepwise procedure (depending on the initial model); `direction='forward'` corresponds to forward selection; `direction='backward'` corresponds to backward elimination.
- The option `scope` specifies the potential pool of X variables (upper) and the X variables that should always be included in the model (lower).
- `k=2` corresponds to *AIC* criterion; `k=log(n)` corresponds to *BIC* criterion.

Surgical Unit: Forward Stepwise

Reading material:

Start with the null-model.

```
> library(MASS)
> fit.0 = lm(log(Y) ~ 1, data=data.o) ## initial model: null-model with only intercept term
> step.0 = stepAIC(fit.0, scope=list(upper="X1+X2+X3+X4+X5+X6+X7+X8", lower="1"), direction="both", k=2)
```

Start: AIC=-75.72

log(Y) ~ 1

	Df	Sum of Sq	RSS	AIC
+ X3	1	5.4708	7.3337	-103.811
+ X4	1	5.3967	7.4079	-103.268
+ X2	1	2.8303	9.9742	-87.205
+ X8	1	1.7808	11.0238	-81.802
+ X1	1	0.7770	12.0275	-77.096
+ X6	1	0.6889	12.1156	-76.703
<none>			12.8045	-75.716
+ X5	1	0.2694	12.5351	-74.864
+ X7	1	0.2067	12.5978	-74.595

Step: AIC=-103.81

log(Y) ~ X3

	Df	Sum of Sq	RSS	AIC
+ X2	1	3.0209	4.3129	-130.479
+ X4	1	2.2018	5.1319	-121.089
+ X1	1	1.5512	5.7825	-114.644
+ X8	1	1.1386	6.1951	-110.922
<none>			7.3337	-103.811
+ X6	1	0.2582	7.0755	-103.747
+ X5	1	0.2390	7.0947	-103.600
+ X7	1	0.0659	7.2679	-102.298
- X3	1	5.4708	12.8045	-75.716

Surgical Unit: Forward Stepwise (Cont'd)

Step: AIC=-130.48

log(Y) ~ X3 + X2

Df	Sum of Sq	RSS	AIC
+ X8	1	1.4709 2.8420	-151.002
+ X1	1	1.2044 3.1085	-146.161
+ X4	1	0.6979 3.6150	-138.011
+ X7	1	0.2280 4.0849	-131.412
+ X5	1	0.1648 4.1481	-130.583
<none>		4.3129	-130.479
+ X6	1	0.0822 4.2306	-129.518
- X2	1	3.0209 7.3337	-103.811
- X3	1	5.6613 9.9742	-87.205

Step: AIC=-151

log(Y) ~ X3 + X2 + X8

Df	Sum of Sq	RSS	AIC
+ X1	1	0.6642 2.1778	-163.376
+ X4	1	0.4658 2.3761	-158.669
+ X6	1	0.1372 2.7048	-151.674
<none>		2.8420	-151.002
+ X5	1	0.0709 2.7711	-150.367
+ X7	1	0.0241 2.8179	-149.462
- X8	1	1.4709 4.3129	-130.479
- X2	1	3.3531 6.1951	-110.922
- X3	1	4.9403 7.7823	-98.605

Surgical Unit: Forward Stepwise (Cont'd)

Step: AIC=-163.38

$\log(Y) \sim X3 + X2 + X8 + X1$

	Df	Sum of Sq	RSS	AIC
+ X6	1	0.0966	2.0812	-163.826
<none>			2.1778	-163.376
+ X5	1	0.0760	2.1018	-163.293
+ X4	1	0.0415	2.1363	-162.415
+ X7	1	0.0224	2.1554	-161.935
- X1	1	0.6642	2.8420	-151.002
- X8	1	0.9307	3.1085	-146.161
- X2	1	2.9891	5.1670	-118.722
- X3	1	5.4459	7.6237	-97.717

Step: AIC=-163.83

$\log(Y) \sim X3 + X2 + X8 + X1 + X6$

	Df	Sum of Sq	RSS	AIC
+ X5	1	0.0769	2.0043	-163.86
<none>			2.0812	-163.83
- X6	1	0.0966	2.1778	-163.38
+ X7	1	0.0219	2.0593	-162.40
+ X4	1	0.0163	2.0649	-162.25
- X1	1	0.6236	2.7048	-151.67
- X8	1	0.9754	3.0567	-145.07
- X2	1	2.8287	4.9099	-119.48
- X3	1	5.0742	7.1554	-99.14

Surgical Unit: Forward Stepwise (Cont'd)

Step: AIC=-163.86

$\log(Y) \sim X3 + X2 + X8 + X1 + X6 + X5$

	Df	Sum of Sq	RSS	AIC
<none>			2.0043	-163.858
- X5	1	0.0769	2.0812	-163.826
- X6	1	0.0975	2.1018	-163.293
+ X7	1	0.0326	1.9718	-162.743
+ X4	1	0.0022	2.0021	-161.919
- X1	1	0.6284	2.6327	-151.133
- X8	1	0.9011	2.9054	-145.810
- X2	1	2.7644	4.7688	-119.052
- X3	1	5.0752	7.0795	-97.716

> stepAIC

Stepwise Model Path

Analysis of Deviance Table

Initial Model:

$\log(Y) \sim 1$

Final Model:

$\log(Y) \sim X3 + X2 + X8 + X1 + X6 + X5$

Step	Df	Deviance	Resid. Df	Resid. Dev	AIC
1			53	12.804509	-75.71608
2 + X3	1	5.47078352	52	7.333726	-103.81102
3 + X2	1	3.02085553	51	4.312870	-130.47855
4 + X8	1	1.47089284	50	2.841977	-151.00214
5 + X1	1	0.66416961	49	2.177808	-163.37593
6 + X6	1	0.09659084	48	2.081217	-163.82569
7 + X5	1	0.07688125	47	2.004335	-163.85826

Surgical Unit: Forward Stepwise (Cont'd)

- The selected model is $X_1, X_2, X_3, X_5, X_6, X_8$ ($p = 7$) with $AIC_p = -163.858$.
- In this case, the forward selection procedure also selects the same model.

```
## forward selection  
> step.0.f=stepAIC(fit.0, scope=list(upper="X1+X2+X3+X4+X5+X6+X7+X8", lower="1"),  
+ direction="forward", k=2)
```

Surgical Unit: Backward Elimination

Start with the full model with all eight predictors.

```
> fit.f = lm(log(Y) ~ ., data=data.o)
> step.b = stepAIC(fit.f, scope = list(upper = ".", lower = "~1"), direction = "backward", k = 2)
Start: AIC=-160.78
log(Y) ~ X1 + X2 + X3 + X4 + X5 + X6 + X7 + X8
```

	Df	Sum of Sq	RSS	AIC
- X4	1	0.00126	1.9718	-162.74
- X7	1	0.03159	2.0021	-161.92
- X5	1	0.07359	2.0441	-160.80
<none>			1.9705	-160.78
- X6	1	0.08403	2.0545	-160.52
- X1	1	0.31845	2.2890	-154.69
- X8	1	0.84489	2.8154	-143.51
- X2	1	2.09285	4.0634	-123.70
- X3	1	2.98863	4.9591	-112.94

Surgical Unit: Backward Elimination (Cont'd)

Step: AIC=-162.74

log(Y) ~ X1 + X2 + X3 + X5 + X6 + X7 + X8

	Df	Sum of Sq	RSS	AIC
- X7	1	0.0326	2.0043	-163.858
<none>			1.9718	-162.743
- X5	1	0.0876	2.0593	-162.396
- X6	1	0.0969	2.0687	-162.152
- X1	1	0.6269	2.5987	-149.835
- X8	1	0.8438	2.8156	-145.506
- X2	1	2.6755	4.6473	-118.446
- X3	1	5.0934	7.0652	-95.825

Step: AIC=-163.86

log(Y) ~ X1 + X2 + X3 + X5 + X6 + X8

	Df	Sum of Sq	RSS	AIC
<none>			2.0043	-163.858
- X5	1	0.0769	2.0812	-163.826
- X6	1	0.0975	2.1018	-163.293
- X1	1	0.6284	2.6327	-151.133
- X8	1	0.9011	2.9054	-145.810
- X2	1	2.7644	4.7688	-119.052
- X3	1	5.0752	7.0795	-97.716

```
## backward stepwise
```

```
> step.bs=stepAIC(fit.f, scope= list(upper=~., lower=~1),  
+ direction="both", k=2)
```

Again the model $X_1, X_2, X_3, X_5, X_6, X_8$ is selected. Backward stepwise also selects the same model.

Stepwise Procedures: Comments

- Forward stepwise procedure often works better than forward selection when there is .
- Backward procedures are not good when the number of potential X variables, $P - 1$, is . Particularly, they are not feasible when $P > n$, since then the full model can not be fitted.
- A potential disadvantage of forward procedures is the MSE and thus the standard errors of the LS estimators tend to be in the initial steps due to

Stepwise Procedures: Comments

- Forward stepwise procedure often works better than forward selection when there is high multicollinearity.
- Backward procedures are not good when the number of potential X variables, $P - 1$, is large. Particularly, they are not feasible when $P > n$, since then the full model can not be fitted.
- A potential disadvantage of forward procedures is the MSE and thus the standard errors of the LS estimators tend to be overestimated in the initial steps due to underfitting since important X variables are likely to be omitted in those steps.

Model Building: Comments

For the sake of interpretability:

- It is often appropriate to select all the indicator variables corresponding to a qualitative variable as a group (i.e., to be in or out of the model simultaneously).
- **Hierarchical principle:** If higher-order terms (e.g., interactions, powers) are selected, it is often appropriate to include the related lower-order terms as well.