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R LAB 6
03/11/2020

TMATH 410 Computer Assignment 6
Lectures 12-13, Chapters 5, 6, 9, 11

Objectives:

1. Practice evaluating collinearity
2. Use R's model selection utility
3. Perform diagnostics on the final model selected

Data for computer assignment

For this assignment we will investigate a data set from a study in 1975-1976 that investigated the risk of hospital acquired infection in the US. The data were published in the American Journal of Epidemiology in 1980, and obtained from Neter et al. The variables in the data set (for this assignment) are: the average length of stay per patient (LenStay), the average age of patients (Age), mean probability of infection (as a percentage; InfectionRisk), average number of beds in hospital (NumBeds), whether the hospital is affiliated with a medical school (MedSchAff; 1=Yes, 2=No), the average number of patients per day (PatientsPerDay), and the number of nurses in the hospital (NumNurse).

C1. (4) Submit the R script you used for this assignment.

Submitted

C2 (2) Create a summary table of the MedSchAff column and report the number of hospitals affiliated with a medical school and the number not affiliated.

```
> summary(scenic.df$MedSchAff)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1.00	2.00	2.00	1.85	2.00	2.00

C3 (3) Create paired scatterplots for each of the quantitative variables. The easiest way to do this is to simply enter: plot(scenic.df). This creates all pairwise scatterplots for all variables in a data frame. Make sure your plotting window is big enough! For this problem it is ok for the graphic to not be of publication-quality. Include the graph in your report and comment on any patterns you see.

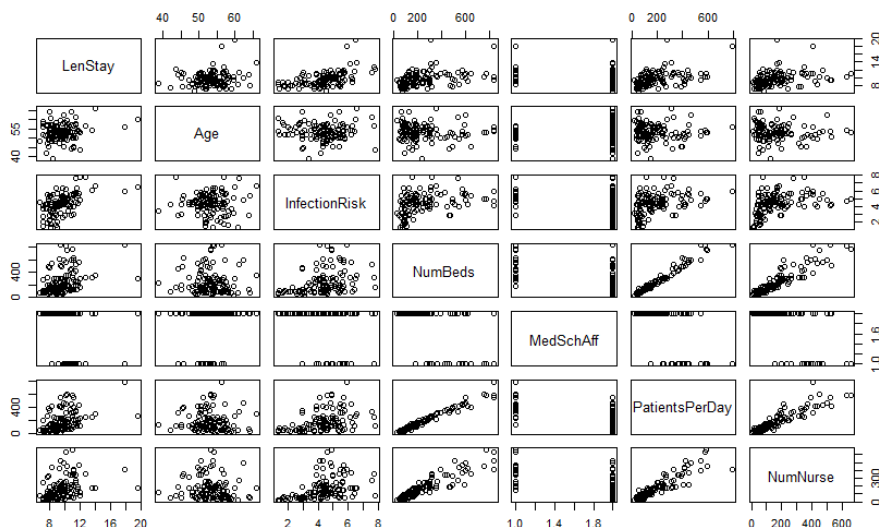


Figure 1 Paired Scatter plot for all quantitative variables

The scatter plots of *MedSchAff* vs. et al are all pointing to the predictor being a factor.

PatientsPerDay correlates positively with **NumNurse**(which also correlates positively with **NumBeds**), as well as with **NumBeds**.

Some of the graphs point to a violation of constant variance, such as infection risk vs each (**NumBeds**, **PatientsPerDay**,**NumNurse**).

C4 (2) Estimate the full model that predicts infection risk as a linear combination of the quantitative explanatory variables. Use the `vif` function in the `car` package to calculate variance inflation factors for the predictors in your linear model, and present the results in a publication-quality table. (Note, you can install the `car` package in RStudio by going to the packages tab, then install, then type `car`. Then type `car::vif(lmObject.lm)` where `lmObject.lm` is the name of your linear model object).

Length of Stay	Age	Number of Beds	Patients per day	Number of Nurse
1.571815	1.071767	31.777783	33.197632	6.450622

Table 1 VIF for the data in the scenic linear model

C5 (2) Comment on the values for VIF, in the context of your graphs in C3.

As can be seen in the linear plots of the predictors with $VIF > 5$, there exists high multicollinearity among these variables which were pointed out in C3.

C6 (2) Estimate a new linear model that predicts InfectionRisk by a linear combination of `LenStay`, `Age`, and `PatientsPerNurse`, then calculate VIFs for this new model and report the results in a publicationquality table. Do these values indicate any issues?

Length of Stay	Age	Patients Per Nurse
1.042785	1.037170	1.006300

Table 2 VIFs after dropping a column and using patients per nurse instead.

These new VIFs do not indicate any underlying issues point to multiple collinearity. They are ideal.

C7. (5) Use the step function model selection for our full linear model using AIC as the criterion. In your lab document summarize which variable was eliminated each step, and explain why.

```
> scenicStep.aic=step(scenic2.lm)
Start: AIC=18.51
InfectionRisk ~ LenStay * MedSchAff + Age * MedSchAff + PatientsPerNurse *
  MedSchAff
```

	Df	Sum of Sq	RSS	AIC
- LenStay:MedSchAff	1	0.29172	115.83	16.795
- MedSchAff:PatientsPerNurse	1	0.44079	115.98	16.941
- MedSchAff:Age	1	1.66883	117.21	18.131
<none>			115.54	18.511

The first parameter to be removed is the *LenStay:MedSchAff* which R removes due to how the interaction between the parameters, once removed, results in a lower AIC = 16.8

```
Step: AIC=16.8
InfectionRisk ~ LenStay + MedSchAff + Age + PatientsPerNurse +
```

MedSchAff:Age + MedSchAff:PatientsPerNurse

	Df	Sum of Sq	RSS	AIC
- MedSchAff:Age	1	1.443	117.27	16.195
- MedSchAff:PatientsPerNurse	1	1.750	117.58	16.490
<none>			115.83	16.795
- LenStay	1	52.352	168.18	56.936

The next step parameter removed is *MedSchAff:Age(interaction)* which R removes due to how the interaction between the parameters, once removed, results in a lower AIC = 16.19

Step: AIC=16.19

InfectionRisk ~ LenStay + MedSchAff + Age + PatientsPerNurse + MedSchAff:PatientsPerNurse

	Df	Sum of Sq	RSS	AIC
- Age	1	1.968	119.24	16.075
<none>			117.27	16.195
- MedSchAff:PatientsPerNurse	1	2.554	119.83	16.630
- LenStay	1	56.846	174.12	58.856

The next step parameter removed is *Age(main)* which R removes due to how the parameters, once removed, results in a lower AIC = 16.08

Step: AIC=16.08

InfectionRisk ~ LenStay + MedSchAff + PatientsPerNurse + MedSchAff:PatientsPerNurse

	Df	Sum of Sq	RSS	AIC
- MedSchAff:PatientsPerNurse	1	2.095	121.34	16.043
<none>			119.24	16.075
- LenStay	1	55.242	174.48	57.092

The next step parameter removed is *MedSchAff:PatientsPerNurse(interaction)* which R removes due to how the interaction between the parameters, once removed, results in a lower AIC = 16.04

Step: AIC=16.04

InfectionRisk ~ LenStay + MedSchAff + PatientsPerNurse

	Df	Sum of Sq	RSS	AIC
- MedSchAff	1	0.726	122.06	14.717
<none>			121.34	16.043
- PatientsPerNurse	1	21.509	142.84	32.484
- LenStay	1	53.179	174.51	55.112

Last step parameter removed is *MedSchAff(main)* which R removes due to how the main effect of the parameter, once removed, results in a lower AIC = 14.717

Step: AIC=14.72

InfectionRisk ~ LenStay + PatientsPerNurse

	Df	Sum of Sq	RSS	AIC
<none>			122.06	14.717
- PatientsPerNurse	1	22.012	144.07	31.453
- LenStay	1	62.624	184.69	59.513

C8. (3) Produce a summary of the final model selected by the step function. Provide an interpretation of each partial regression coefficient

$$\hat{Y} = 1.5741 + 0.3924\hat{X}_1\beta_1 - 0.8450\hat{X}_2\beta_2$$

```
> scenicStep.aic
```

```
Call:
```

```
lm(formula = InfectionRisk ~ LenStay + PatientsPerNurse, data = scenic.df)
```

```
Coefficients:
```

```
(Intercept)          LenStay  PatientsPerNurse
```

```
> summary(scenicStep.aic)
```

```
Call:
```

```
lm(formula = InfectionRisk ~ LenStay + PatientsPerNurse, data = scenic.df)
```

```
Residuals:
```

```
      Min       1Q   Median       3Q      Max
-2.6062  -0.5384  -0.0393   0.6126   2.7684
```

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.57407	0.54494	2.889	0.00466
LenStay	0.39240	0.05223	7.512	1.64e-11
PatientsPerNurse	-0.84504	0.18973	-4.454	2.03e-05

```
(Intercept)      **
```

```
LenStay          ***
```

```
PatientsPerNurse ***
```

```
---
```

```
Signif. codes:
```

```
0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Residual standard error: 1.053 on 110 degrees of freedom
```

```
Multiple R-squared:  0.3939, Adjusted R-squared:  0.3828
```

```
F-statistic: 35.74 on 2 and 110 DF, p-value: 1.099e-12
```

Interpretation: We see that for every unit increase in Length of Patient Stay, after effects of Patients Per Nurse are accounted for, the infection risk goes up by a factor of 0.39240 plus a standard 1.57407units. Additionally, after the effects of Length of stay are accounted for, the infection risk decreases by a factor of 0.84504 plus the standard 1.57407, for every unit increase in Patients Per Nurse.

C9. (3) Give residual plots of your final model. Comment on any patterns you see, and whether you believe any major regression assumptions are violated. Decide if there are any influential points in your model.

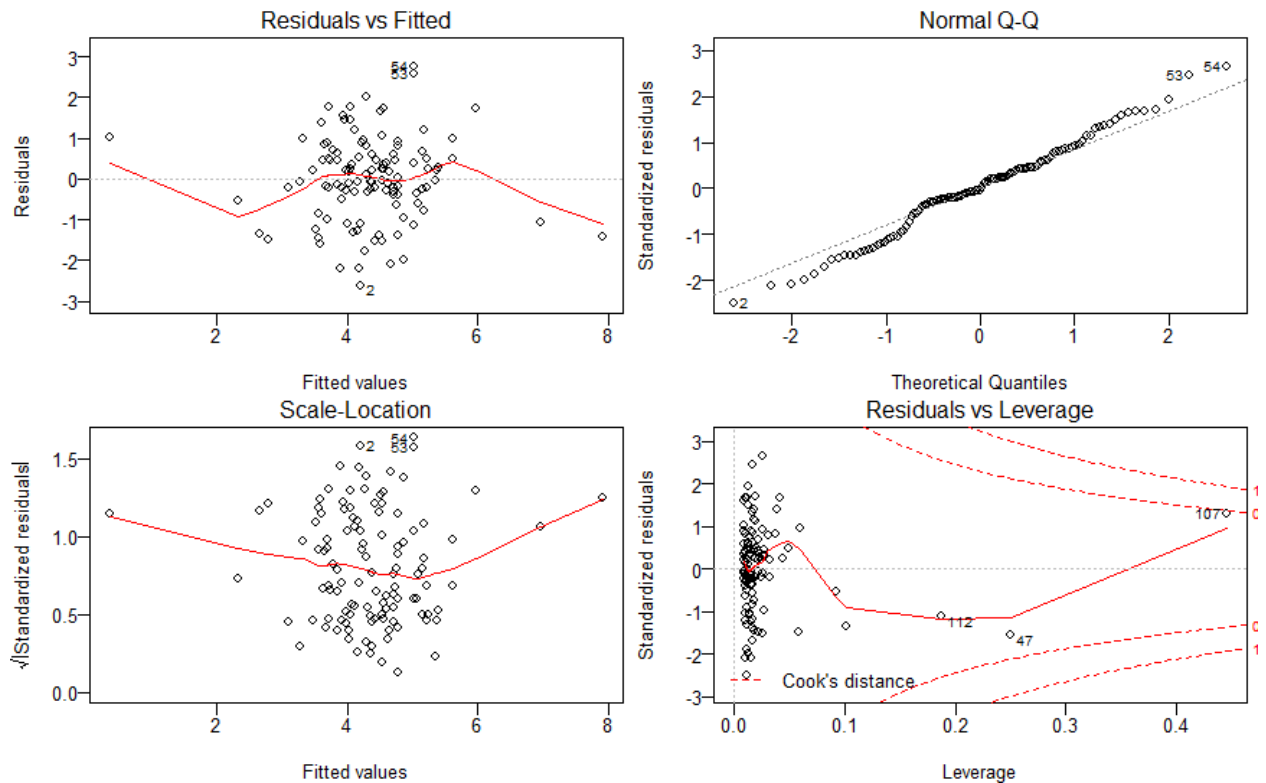


Table 3 Residual diagnosis plots generated using R

The cook's plot as can be seen in the fourth plot, does not highlight points that might be problematic and of great influence on our model. Instead, the other three plots highlight outlier points 53 and 54, which also seem to have high leverage. Outside from these two points, the plots are pointing to a residual distribution that would be decent since constant variance assumption is retained and the residual points are independently and identically distributed.

C10 (4) Interpret the results of your model selection procedure in the context of what we have learned about infection risk through this data analysis. Include in your discussion consideration of which variables were included in the final model and which variables were not included.

Even though there is a high correlation among some variables, the VIF pointed out multicollinearity in the data set, which was at first hinted to by the initial plots. This confirmation enabled the points that had significant VIFs to be considered and removed or used to estimate a new model with an additional column that was the Patients Per Nurse. The model selection using the AIC criterion facilitated the arrival of a model that was simple and fit the data appropriately, which was limited to just two predictor variables which had the lowest AIC score of 14.72 (without any additional interacting predictors). But in sum, we found that the infection risk could be predicted or explained by taking into account the number of patients per nurse and the length/number of days patient stayed in the hospital; and this infection risk was inversely proportional to the former, but directly proportional to the latter.