

# STAT 408

# Applied Regression Analysis

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# Transformation

# Transformation

- In linear regression, the “linear” refers to parameter
- The predictors themselves do not have to be linear

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 \log X_2 + \beta_3 X_1 X_2 + \varepsilon$$



$$Y = \beta_0 + \beta_1 X_1^{\beta_2} + \varepsilon$$



- Therefore, we can transform both response Y and predictors and model is still linear
- Let's first look at transforming response Y

# Transforming the Response

- Reasons we may consider transforming response  $Y$  in linear model
  1. Reduce the impact of outliers and increase the normality of error distribution
  2. Improve the model fit
  3. Some real questions require us to transform the response  $Y$
- We already see log and square root transformation for 1 and 2
- Let's see one example for 3

# Transforming the Response

- Production function in microeconomics shows

$$Y = AL^{\beta}K^{\alpha}$$

where  $Y$  = total production,  $L$  = labor input,  $K$  = capital input,  $A$  = productivity  
 $\alpha$  and  $\beta$  are “elasticity” of labor and capital, which are our interests

- Log-transformation gives us a linear model to estimate  $\alpha$  and  $\beta$ :

$$\log(Y) = \log(A) + \beta \log(L) + \alpha \log(K)$$

# Transforming the Response

- When we use log-transformation, the regression parameters have a particular interpretation

$$\hat{y} = e^{\hat{\beta}_0} e^{\hat{\beta}_1 x_1} \dots e^{\hat{\beta}_p x_p}$$

$$\log \hat{y} = \hat{\beta}_0 + \hat{\beta}_1 x_1 + \dots + \hat{\beta}_p x_p$$

- If  $\hat{\beta}_1 = 0.01$ , then one unit increase of  $X_1$  will increase  $\log(\hat{y})$  by 0.01
- Because  $\log(1 + \hat{y} - 1) = 0.01$  and  $\log(1+x) \approx x$  for small  $x$ ,  $\hat{y} - 1 \approx 0.01$ 
  - $\hat{y}$  will increase from 1 to 1.01, that is, 1%
- After log-transformation, small  $\hat{\beta}$  is the percentage increase of  $Y$  if  $X$  increases by one unit

# Segmented Regression

- Now let's focus on the transformation of predictors – segmented regression
- The saving dataset contains five variables for 50 countries
  - sr: saving rate
  - pop15: percent population under age of 15
  - pop75: percent population over age of 75
  - dpi: per-capita disposable income in dollars
  - ddpi: percent growth rate of dpi
- The data is over the period 1960-1970

	sr	pop15	pop75	dpi	ddpi
<b>Australia</b>	11.43	29.35	2.87	2329.68	2.87
<b>Austria</b>	12.07	23.32	4.41	1507.99	3.93
<b>Belgium</b>	13.17	23.80	4.43	2108.47	3.82
<b>Bolivia</b>	5.75	41.89	1.67	189.13	0.22
<b>Brazil</b>	12.88	42.19	0.83	728.47	4.56
<b>Canada</b>	8.79	31.72	2.85	2982.88	2.43

# Segmented Regression

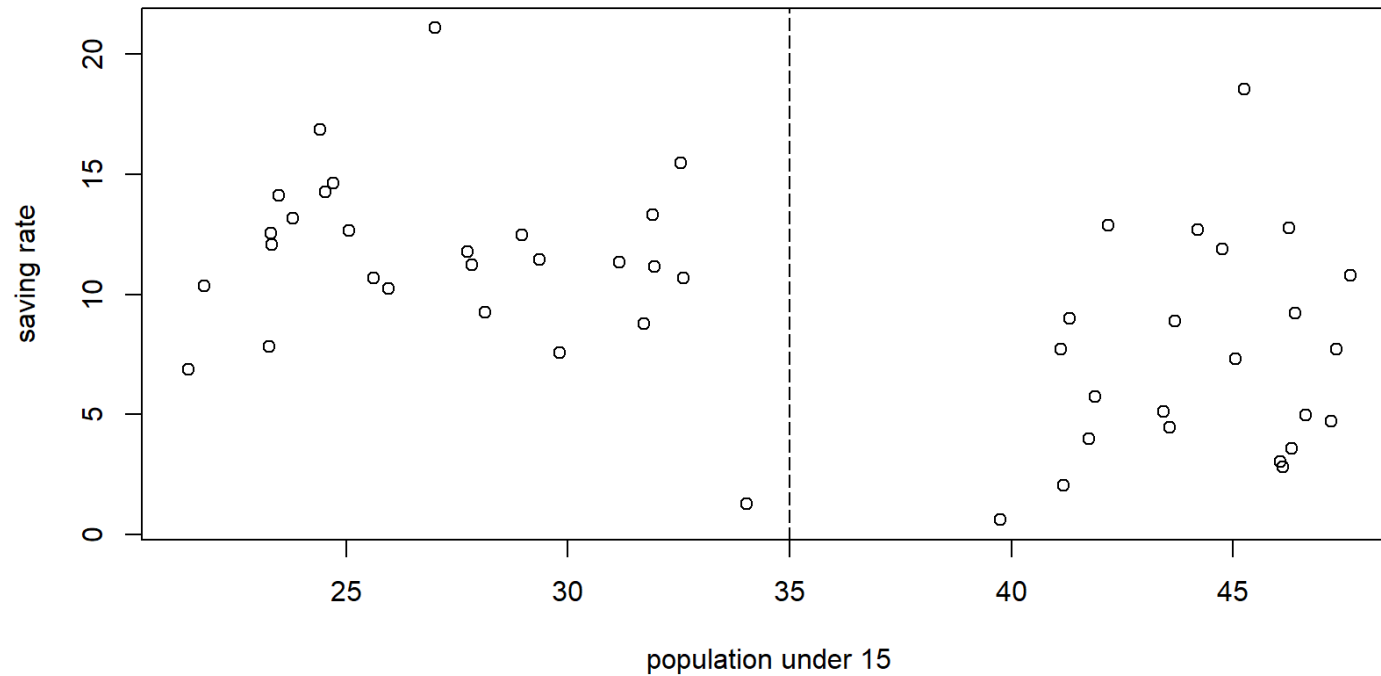
- The motivation of segmented regression is that different linear regression models may apply in different regions of the data
- In the saving dataset, we suspect the relations between saving rate and population age are different in younger countries and older countries
- We use  $\text{pop15}=35$  as the cutoff

```
saving <- read.csv("saving.csv")  
plot(sr ~ pop15, data = saving, xlab='population under 15', ylab='saving rate')  
abline(v=35, lty=5)
```



# Segmented Regression

- We need to fit two separate models to capture the two relationships



# Segmented Regression

- Fit two models in two regions

```
lm1 <- lm(sr~pop15, data = saving, subset = (pop15<35))
```

```
lm2 <- lm(sr~pop15, data = saving, subset = (pop15>35))
```

- Draw two models in two regions

```
segments(x0 = 20, y0 = lm1$coefficients[1]+lm1$coefficients[2]*20,
```

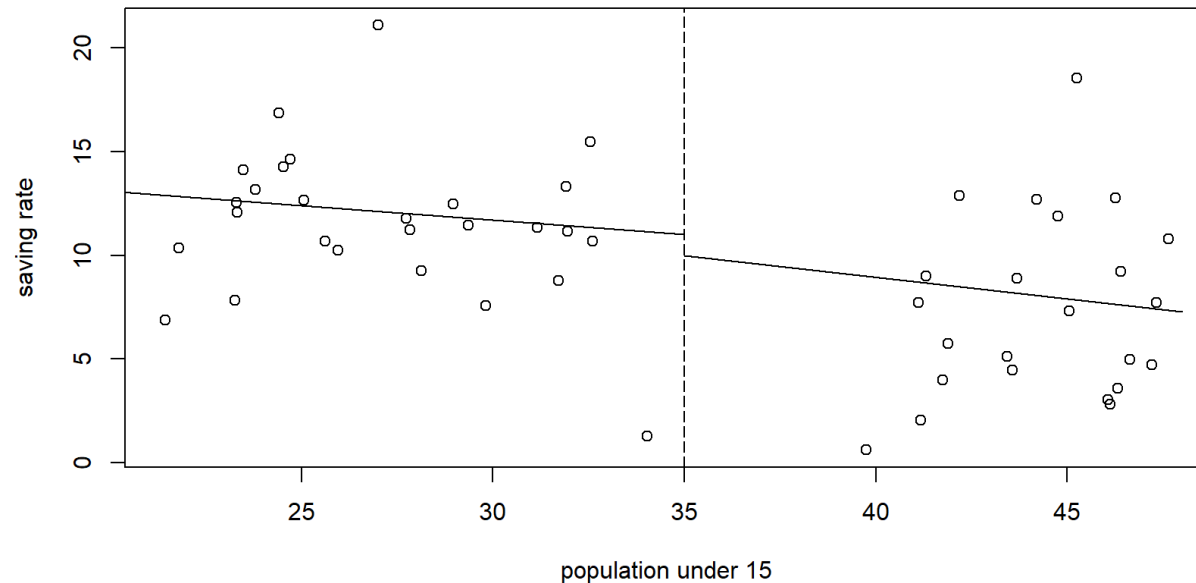
```
        x1 = 35, y1 = lm1$coefficients[1]+lm1$coefficients[2]*30)
```

```
segments(x0 = 35, y0 = lm2$coefficients[1]+lm2$coefficients[2]*35,
```

```
        x1 = 48, y1 = lm2$coefficients[1]+lm2$coefficients[2]*48)
```

# Segmented Regression

- One issue is that the two models are disconnected at  $x=35$
- $X=35$  is a break point, which seems unrealistic
- Segmented regression solves this issue by smoothing the break points



# Segmented Regression

- We transform the predictor  $X$  into two predictors by two basis functions

$$B_l(x) = \begin{cases} c - x & \text{if } x < c \\ 0 & \text{otherwise} \end{cases}$$

$$B_r(x) = \begin{cases} x - c & \text{if } x > c \\ 0 & \text{otherwise} \end{cases}$$

where  $c$  is the cutoff between the two groups and  $c = 35$  in this model

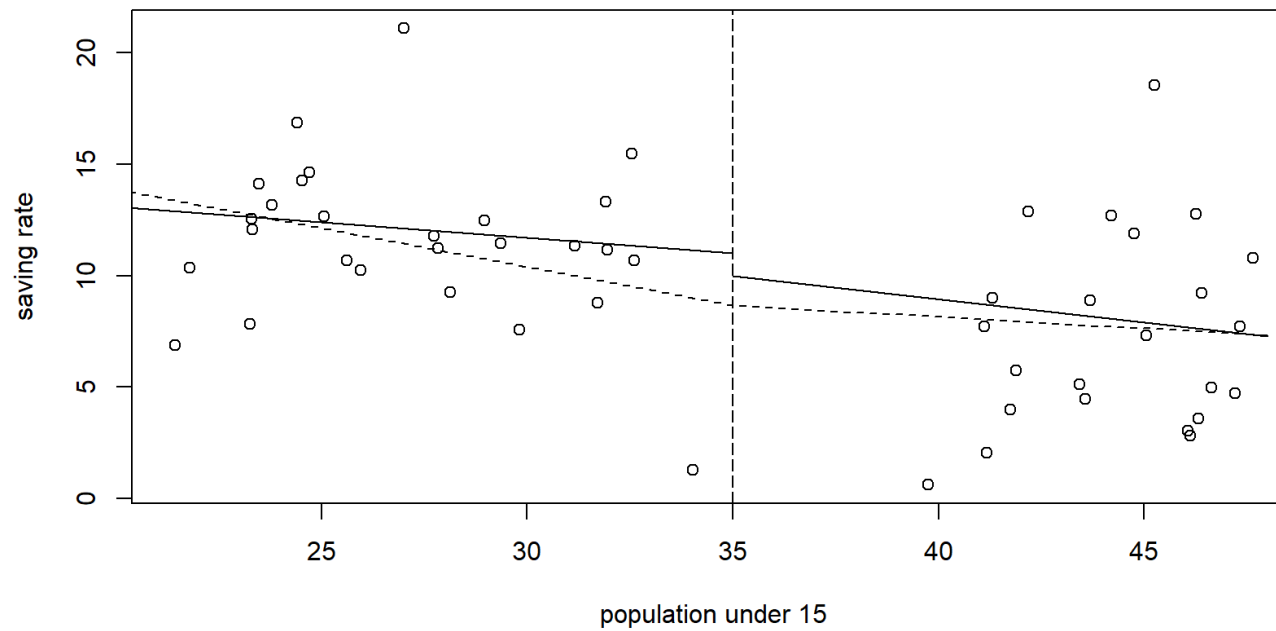
- Under this transformation, the linear model is

$$y = \beta_0 + \beta_1 B_l(x) + \beta_2 B_r(x) + \varepsilon$$

- We can use regular linear regression to fit this model ([coding 6.r](#))

# Segmented Regression

- The segmented model will have different slopes in the two groups but connects at  $X = 35$
- Segmented model also reduces the four parameters in the two linear models to three
- Since we change the  $X$  value in both groups, the two slopes are different from the two separate regressions



# Polynomial Regression

- Polynomial regression includes the higher order of predictors into a linear model

$$y = \beta_0 + \beta_1 x + \cdots + \beta_d x^d + \varepsilon$$

- It introduces non-linearity into the model and provides more flexibility and complexity
- There are three ways to determine the higher order  $d$ 
  1. Forward: adding terms until the added term is not statistically significant
  2. Backward: starting with a large  $d$  and eliminate non-statistically significant terms
  3. Choose  $d$  based on prior knowledge

# Polynomial Regression

- We use forward method to examine the relation between saving and the increase of disposable income

```
> summary(lm(sr ~ ddpi, savings))
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	7.883	1.011	7.80	4.5e-10
ddpi	0.476	0.215	2.22	0.031

```
> summary(lm(sr ~ ddpi+I(ddpi^2), savings))
```

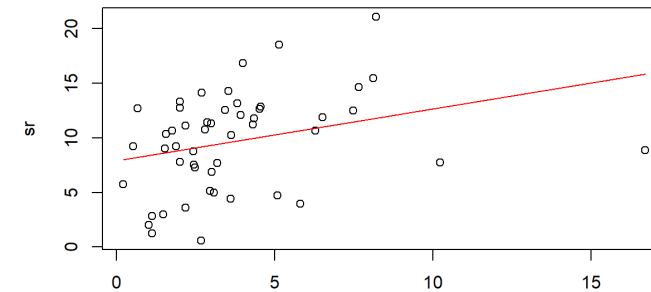
Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	5.1304	1.4347	3.58	0.00082
ddpi	1.7575	0.5377	3.27	0.00203
I(ddpi^2)	-0.0930	0.0361	-2.57	0.01326

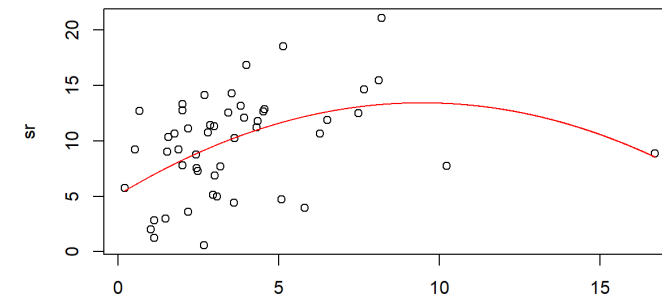
```
> summary(lm(sr ~ ddpi+I(ddpi^2)+I(ddpi^3), savings))
```

Coefficients:

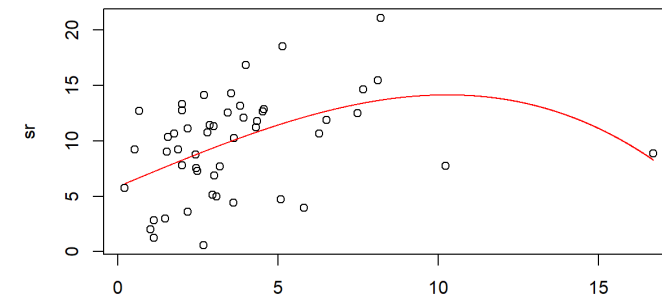
	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	5.145360	2.198606	2.34	0.024
ddpi	1.746017	1.380455	1.26	0.212
I(ddpi^2)	-0.090967	0.225598	-0.40	0.689
I(ddpi^3)	-0.000085	0.009374	-0.01	0.993



Linear



Quadratic



Cubic

# Polynomial Regression

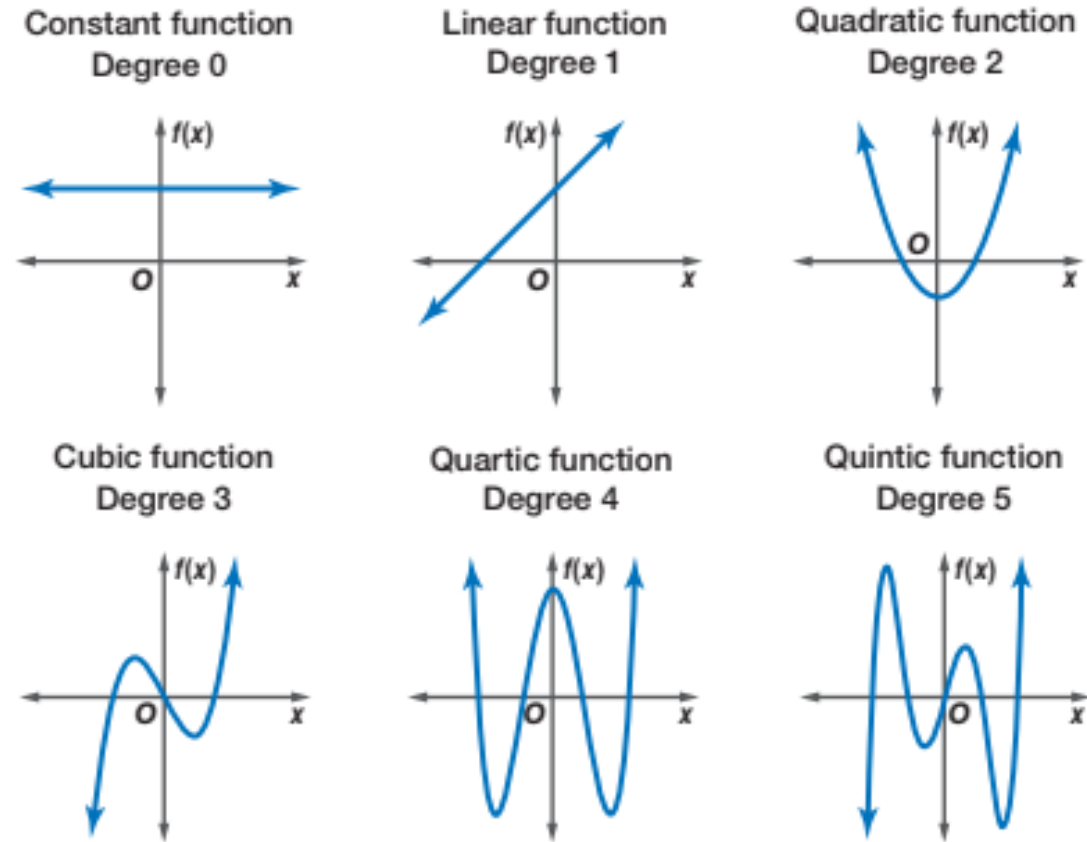
- Any term higher than the second order is insignificant, so the final model is quadratic of dppi
- It is a bad idea to eliminate lower order terms from the model before the higher order terms, even if they are not significant
- We can also define polynomials in more than one variable, also called response surface model

$$y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_{11} x_1^2 + \beta_{22} x_2^2 + \beta_{12} x_1 x_2$$

```
lmod <- lm(sr ~ pop15 + ddpi + I(pop15^2) + I(ddpi^2) + I(pop15*ddpi), savings)
summary(lmod)
```



# Nonlinearity in Linear Regression



The visualization for polynomial models with different orders

# Collinearity

# Collinearity

- Recall that we estimate linear model by

$$\hat{\beta} = (X^T X)^{-1} X^T y$$

- If  $X$  is singular (perfect linear relation of predictors), then  $X^T X$  is not invertible,  $\hat{\beta}$  does not have unique solution
- In this case, we need to drop certain predictors to break perfect linear relation
- This is called exact collinearity

# Collinearity

- A more challenging problem is  $X$  close to singular but not exactly (collinearity)
- Recall  $\hat{\beta}$  is a random variable with a normal distribution:

$$\hat{\beta} \sim N(\beta, (X^T X)^{-1} \sigma^2)$$

- Close to singular will cause  $X^T X$  “small” and  $(X^T X)^{-1}$  “large”, then the variance of  $\hat{\beta}$  will be large
  1. The model estimation is unstable, small measurement errors leads to large changes in  $\hat{\beta}$
  2. t statistic is small, t-test may fail to find significant predictors
  3. The signs of the coefficients could be the opposite of the truth

# Collinearity Detection

1. Examine the correlation matrix of the predictors
  - Matrix entry close to  $-1$  or  $+1$  indicates large pairwise collinearities
2. Regress predictor  $X_i$  on all other predictors, then check R-square of this regression
  - large R-square (close to one) indicates collinearity

# Collinearity Detection

- Let's see one real-data example of dealing with collinearity
- Dataset seatpos contains 8 predictors of 38 driver's body size, weight, age and response variable hipcenter (seating position)

	Age	Weight	HtShoes	Ht	Seated	Arm	Thigh	Leg	hipcenter
1	46	180	187.2	184.9	95.2	36.1	45.3	41.3	-206.300
2	31	175	167.5	165.5	83.8	32.9	36.5	35.9	-178.210
3	23	100	153.6	152.2	82.9	26.0	36.6	31.0	-71.673
4	19	185	190.3	187.4	97.3	37.4	44.1	41.0	-257.720
5	23	159	178.0	174.1	93.9	29.5	40.1	36.9	-173.230
6	47	170	178.7	177.0	92.4	36.0	43.2	37.4	-185.150
7	30	137	165.7	164.6	87.7	32.5	35.6	36.2	-164.750
8	28	192	185.3	182.7	96.9	35.8	39.9	43.1	-270.920

# Collinearity Detection

```
> lmod <- lm(hipcenter ~ ., seatpos)
              Estimate Std. Error t value Pr(>|t|)
(Intercept) 436.43213   166.57162    2.620  0.0138 *
Age          0.77572     0.57033     1.360  0.1843
Weight       0.02631     0.33097     0.080  0.9372
HtShoes     -2.69241     9.75304    -0.276  0.7845
Ht           0.60134    10.12987     0.059  0.9531
Seated       0.53375     3.76189     0.142  0.8882
Arm         -1.32807     3.90020    -0.341  0.7359
Thigh       -1.14312     2.66002    -0.430  0.6706
Leg         -6.43905     4.71386    -1.366  0.1824
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 37.72 on 29 degrees of freedom
Multiple R-squared:  0.6866,    Adjusted R-squared:  0.6001
F-statistic:  7.94 on 8 and 29 DF,  p-value: 1.306e-05
```

## Issue of this full model

- No single predictor is significant (t-test), but they are jointly significant (F-test)
- Standard deviations of estimated parameters are large
- The R-square is reasonable
- Multiple predictors measure driver's body size
- Those are evidence of collinearity: highly correlated predictors

# Collinearity Detection

- Let's check the predictor correlation matrix

```
> round(cor(seatpos[, -9]), 2)
```

	Age	Weight	HtShoes	Ht	Seated	Arm	Thigh	Leg
Age	1.00	0.08	-0.08	-0.09	-0.17	0.36	0.09	-0.04
Weight	0.08	1.00	0.83	0.83	0.78	0.70	0.57	0.78
HtShoes	-0.08	0.83	1.00	1.00	0.93	0.75	0.72	0.91
Ht	-0.09	0.83	1.00	1.00	0.93	0.75	0.73	0.91
Seated	-0.17	0.78	0.93	0.93	1.00	0.63	0.61	0.81
Arm	0.36	0.70	0.75	0.75	0.63	1.00	0.67	0.75
Thigh	0.09	0.57	0.72	0.73	0.61	0.67	1.00	0.65
Leg	-0.04	0.78	0.91	0.91	0.81	0.75	0.65	1.00

- There are some large pairwise correlations between predictors, mainly those predictors that measure height/length



# Collinearity Detection

- Let's regress each predictor on others and check their R squares

```
x <- model.matrix(lmod)[,-1]

for(i in 1:8){
  r2 <- summary(lm(x[,i] ~ x[,-i]))$r.squared
  cat(colnames(x)[i], '\t', r2, '\n')
}
```

Age	0.4994823
Weight	0.7258043
HtShoes	0.9967472
Ht	0.9969982
Seated	0.8882813
Arm	0.7775983
Thigh	0.6380596
Leg	0.850619

- There are some large R-squares indicating collinearity

# Unstable estimation

- We simulate a new dataset by adding random noise (std = 10) to response variable hipcenter

```
lm.model <- lm(hipcenter+rnorm(n=38,mean=0,sd=10)~., data = seatpos)
summary(lm.model)
```

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	420.9929	172.4276	2.442	0.021 *
Age	0.7916	0.5904	1.341	0.190
Weight	-0.1128	0.3426	-0.329	0.744
HtShoes	-6.2007	10.0959	-0.614	0.544
Ht	4.1962	10.4860	0.400	0.692
Seated	0.3776	3.8941	0.097	0.923
Arm	-0.6793	4.0373	-0.168	0.868
Thigh	-1.1333	2.7535	-0.412	0.684
Leg	-5.8780	4.8796	-1.205	0.238

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	436.43213	166.57162	2.620	0.0138 *
Age	0.77572	0.57033	1.360	0.1843
Weight	0.02631	0.33097	0.080	0.9372
HtShoes	-2.69241	9.75304	-0.276	0.7845
Ht	0.60134	10.12987	0.059	0.9531
Seated	0.53375	3.76189	0.142	0.8882
Arm	-1.32807	3.90020	-0.341	0.7359
Thigh	-1.14312	2.66002	-0.430	0.6706
Leg	-6.43905	4.71386	-1.366	0.1824

- Many “length-related” predictors have very different parameter estimations, some even change signs

# Mitigation of Collinearity

- Too many variables try do the same job of explaining the response and there is redundant information in predictors
- When we have a new dataset from the same population, the model “randomly” reassign importance to similar predictors and causes instable parameter estimation
- The high degree of instability inflates the variance of estimation and hides the significance
- The solution is simple: remove highly correlated predictors, leave remain only one of them

# Mitigation of Collinearity

- Let's remove all “length-related” predictors except driver's height

```
lm.model <- lm(hipcenter~Age+Weight+Ht, data = seatpos)
summary(lm.model)
```

```
Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept)  528.297729  135.312947   3.904 0.000426 ***
Age           0.519504   0.408039   1.273 0.211593
Weight       0.004271   0.311720   0.014 0.989149
Ht          -4.211905   0.999056  -4.216 0.000174 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 36.49 on 34 degrees of freedom
Multiple R-squared:  0.6562,    Adjusted R-squared:  0.6258
F-statistic: 21.63 on 3 and 34 DF,  p-value: 5.125e-08
```

- In the new model, the standard deviations of estimated parameters is much smaller
- The height predictor now is significant
- Adjusted  $R^2$  is improved