



ST518 - Multiple linear regression module

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Multiple linear regression(MLR)

Toy example: A random sample of students taking the same exam:

| IQ | Study TIME | GRADE |
|-----|------------|-------|
| 105 | 10 | 75 |
| 110 | 12 | 79 |
| 120 | 6 | 68 |
| 116 | 13 | 85 |
| 122 | 16 | 91 |
| 130 | 8 | 79 |
| 114 | 20 | 98 |
| 102 | 15 | 76 |

Consider a regression model for the GRADE of subject i , Y_i , in which the mean of Y_i is a linear function of two predictor variables $X_{i1} = \text{IQ}$ and $X_{i2} = \text{Study TIME}$ for subjects $i = 1, \dots, 8$:

$$Y = \beta_0 + \beta_1 \text{IQ} + \beta_2 \text{TIME} + \text{error}$$

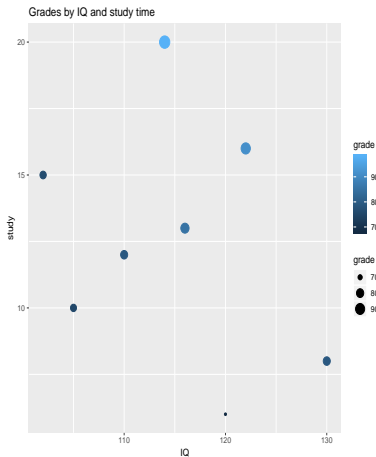
or

$$Y_i = \beta_0 + \beta_1 X_{i1} + \beta_2 X_{i2} + E_i \quad \text{for } i = 1, \dots, 8$$

```

> library(ggplot2)
> iqstudy.plot <- ggplot(iqstudy.dat, aes(IQ, study))
> iqstudy.plot + geom_point(aes(color=grade, size=grade))
+ ggtitle("Grades by IQ and study time")

```



MLR model w/ p independent variables

- Observed values of p independent/predictor variables for i^{th} subject from sample denoted by $x_i. = (x_{i1}, x_{i2}, \dots, x_{ip})$
- response variable for i^{th} subject denoted by Y_i
- For $i = 1, \dots, n$, MLR model for Y_i :

$$Y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots + \beta_p x_{ip} + E_i.$$

- As in SLR, $E_1, \dots, E_n \stackrel{iid}{\sim} N(0, \sigma^2)$, or at least $IND(0, \sigma^2)$.

Least squares estimates of regression parameters (β_i) minimize $SS[E]$:

$$SS[E] = \sum_{i=1}^n (y_i - (\beta_0 + \beta_1 x_{i1} + \dots + \beta_p x_{ip}))^2$$

$$\hat{\sigma}^2 = \frac{SS[E]}{n-p-1}$$

Interpretations of regression parameters:

- σ^2 is unknown error variance parameter.
- $\beta_0, \beta_1, \dots, \beta_p$ are $p + 1$ unknown regression parameters:
 - β_0 : average response when $x_1 = x_2 = \dots = x_p = 0$
 - β_i is called a slope for x_i . Represents mean change in y per unit increase in x_i **with all other independent variables held fixed**.

For the IQ/study time example, with $p = 2$ and $n = 8$,

$$\hat{\beta}_0 = 0.74, \hat{\beta}_1 = 0.47, \hat{\beta}_2 = 2.1$$

What is the uncertainty associated with these parameter estimates?

Is $\beta_1 = 0$ and/or $\beta_2 = 0$ consistent with data?

(Statistical inference.)

Matrix formulation of MLR

Let $x_{i\cdot}$ be a row vector for p observed independent variables for individual i

$$x_{i\cdot} = (1, x_{i1}, x_{i2}, x_{i3}, \dots, x_{ip}) \quad (1 \times (p+1)).$$

MLR model for Y_1, \dots, Y_n given by

$$\begin{aligned} Y_1 &= \beta_0 + \beta_1 x_{11} + \beta_2 x_{12} + \dots + \beta_p x_{1p} + E_1 \\ Y_2 &= \text{_____} \quad (\text{fill this in!}) \\ \vdots &= \vdots \\ Y_n &= \beta_0 + \beta_1 x_{n1} + \beta_2 x_{n2} + \dots + \beta_p x_{np} + E_n \end{aligned}$$

System of n equations can be expressed using matrices: $Y = X\beta + E$

- Y denotes a response vector ($n \times 1$)
- X denotes a _____ matrix ($n \times (p+1)$)
- β denotes a vector of regression parameters ($(p+1) \times 1$)
- E denotes an error vector ($n \times 1$), assumed $MVN(0, \sigma^2 I_n)$.

To obtain matrix expressions for the LS estimates of β , take partial derivatives of the sum of squares function,

$$\begin{aligned} Q(\beta) &= \sum Y_i - (\beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \cdots + \beta_p x_{ip})^2 \\ &= (Y - X\beta)'(Y - X\beta) \\ &= \end{aligned}$$

Note that if b is $p \times 1$ and A is $(p \times p)$, then $\frac{\partial b'Ab}{\partial b} = (A + A')b$.

$$\frac{\partial Q}{\partial \beta} = -2X'Y + (X'X + (X'X)')\beta$$

The $p + 1$ equations with $p + 1$ unknowns obtained by setting this vector of partial derivatives are called the **normal equations**.

$$\frac{\partial Q}{\partial \beta} = 0 \implies \hat{\beta} = \end{aligned}$$

Moments of linear combinations of random vectors

Let Y denote a $p \times 1$ random vector with mean μ and covariance matrix Σ . Suppose a is a $p \times 1$ (fixed) vector of coefficients. Then

$$\begin{aligned} E(a'Y) &= a'\mu \\ \text{Var}(a'Y) &= a'\Sigma a. \end{aligned}$$

So, let's derive $\text{Var}(\hat{\beta}|X)$:

$$\begin{aligned} \text{Var}(\hat{\beta}|X) &= V[(X'X)^{-1}X'Y|X] \\ &= ((X'X)^{-1}X')\text{Var}(Y|X)((X'X)^{-1}X')' \\ &= ((X'X)^{-1}X')\sigma^2 I_n((X'X)^{-1}X')' \\ &= \sigma^2((X'X)^{-1}X')((X'X)^{-1}X')' \\ &= \sigma^2(X'X)^{-1}X'X(X'X)^{-1} \text{(untranspose RHS)} \\ &= \sigma^2 \underline{\hspace{10cm}} \end{aligned}$$

The variance-covariance matrix of the estimated regression coefficients.

$$\begin{aligned}
 \hat{\beta} &= (X'X)^{-1}X'Y \\
 \text{Var}(\hat{\beta}) &= \sigma^2(X'X)^{-1} \\
 &= \Sigma \\
 \widehat{\text{Var}}(\hat{\beta}) &= MS[E](X'X)^{-1} \\
 &= \hat{\Sigma} \\
 \widehat{\text{Var}}(a'\hat{\beta}) &= a'\hat{\Sigma}a
 \end{aligned}$$

- $(X'X)^{-1}$ verbalized as “x prime x inverse”
- X assumed to be of full *rank*

$$\begin{aligned}
 \hat{Y} &= X\hat{\beta} = X(X'X)^{-1}X'Y = HY \\
 e &= Y - \hat{Y} = Y - X\hat{\beta} = (I - H)Y \\
 e'e &= (Y - \hat{Y})'(Y - \hat{Y}) = Y'(I - H)'(I - H)Y = Y'(I - H)Y
 \end{aligned}$$

- \hat{Y} is called the vector of fitted or predicted values
- $H = X(X'X)^{-1}X'$ is called the hat matrix. It is *idempotent*.
- e is the vector of residuals

IQ, Study TIME example, $p = 2$ predictors and $n = 8$ observations, consider $X, Y, (X'X)^{-1}, (X'X)^{-1}X; Y, X(X'X)^{-1}X'Y$:

$$X = \begin{pmatrix} 1 & 105 & 10 \\ 1 & 110 & 12 \\ 1 & 120 & 6 \\ 1 & 116 & 13 \\ 1 & 122 & 16 \\ 1 & 130 & 8 \\ 1 & 114 & 20 \\ 1 & 102 & 15 \end{pmatrix}, \quad X'X = \begin{pmatrix} 8 & 919 & 100 \\ 919 & 106165 & 11400 \\ 100 & 11400 & 1394 \end{pmatrix}$$

$$(X'X)^{-1} = \begin{pmatrix} 28.90 & -0.23 & -0.22 \\ -0.23 & 0.0018 & 0.0011 \\ -0.22 & 0.0011 & 0.0076 \end{pmatrix}$$

$$(X'X)^{-1}X'Y = \begin{pmatrix} 0.74 \\ 0.47 \\ 2.10 \end{pmatrix} = ?$$

$$SS[E] = e'e = (Y - \hat{Y})'(Y - \hat{Y}) = 45.8, \quad e'e/df = 9.15 = ?$$

$$\hat{\Sigma} = MS[E](X'X)^{-1} = \begin{pmatrix} 264.45 & -2.07 & -2.05 \\ -2.07 & 0.017 & 0.010 \\ -2.05 & 0.010 & 0.070 \end{pmatrix}$$

Distribution of parameter estimators,

- If $E \sim N(0, \sigma^2 I)$, then the LS estimator, $\hat{\beta} \sim N(\beta, \Sigma)$. and
- the t -statistics formed from $t = (\hat{\beta}_j - \beta_j) / \sqrt{\widehat{\Sigma}_{jj}}$ follow t -distributions with $df = n - p - 1$.
- If $E \sim IND(0, \sigma^2)$, the normality of $\hat{\beta}$ is approximate.

Some questions - use preceding pages

- ① What is the estimate for β_1 ? Interpretation?
- ② What is the standard error of $\hat{\beta}_1$?
- ③ Is $\beta_1 = 0$ plausible, while controlling for possible linear associations between Test Score and Study time? ($t(0.025, 5) = 2.57$)
- ④ Estimate the mean grade among the population of ALL students with $IQ = 113$ who study $TIME = 14$ hours.
- ⑤ Report a standard error for the estimate in ④
- ⑥ Report a 95% confidence interval for the quantity being estimated in ④
- ⑦ Report a 95% prediction interval for an individual student with $IQ = 113$, $TIME = 14$.
- ⑧ Estimate the std. deviation among students whose mean estimated in ④

Some answers

- 1 $\hat{\beta}_1 = 0.47$ (second element of $(X'X)^{-1}X'Y$, exam points per IQ point for students studying the same amount)
- 2 $\sqrt{0.017} = 0.13$ (square root of middle element of $\hat{\Sigma}$)
- 3 $H_0 : \beta_1 = 0$, T-statistic: $t = (\hat{\beta}_1 - 0)/SE(\hat{\beta}_1)$
Observed value is $t = .47/\sqrt{.017} = .47/.13 = 3.6 > 2.57$,
(" $\hat{\beta}_1$ differs significantly from 0.")
- 4 Unknown population mean: $\theta = \beta_0 + \beta_1(113) + \beta_2(14)$
Estimate : $\hat{\theta} = (1, 113, 14) * \hat{\beta} = 83.6$
- 5 $\text{Var}((1, 113, 14) * \hat{\beta}) = (1, 113, 14)\widehat{\text{Var}}(\hat{\beta})(1, 113, 14)'$
or $(1, 113, 14)\hat{\Sigma}(1, 113, 14)' = 1.3$ or $SE(\hat{\theta}) = \sqrt{1.3} = 1.14$ (on $df = \underline{\hspace{2cm}}$)
- 6 $\hat{\theta} \pm t(0.025, 5)SE(\hat{\theta})$ or $83.6 \pm 2.57(1.14)$ or $(80.7, 86.6)$
- 7 $\hat{Y} \pm t(0.025, 5)\sqrt{MS(E) + SE(\hat{\theta})^2}$ or $83.6 \pm 2.57\sqrt{(9.15 + 1.14^2)}$ or $(75.3, 91.9)$
- 8 $\sqrt{MS(E)} = \sqrt{9.15} = 3.0$ (points)

```

DATA GRADES; INPUT IQ STUDY GRADE @@; CARDS;
105 10 75 110 12 79 120 6 68 116 13 85 122 16 91 130 8 79 114 20 98 102 15 76
DATA EXTRA; INPUT IQ STUDY GRADE; CARDS;
113 14 .
DATA BOTH; SET GRADES EXTRA;
PROC REG; MODEL GRADE = IQ STUDY/P CLM XPX INV COVB;

```

The SAS System
The REG Procedure

Model Crossproducts X'X X'Y Y'Y

| Variable | Intercept | IQ | STUDY | GRADE |
|-----------|-----------|--------|-------|-------|
| Intercept | 8 | 919 | 100 | 651 |
| IQ | 919 | 106165 | 11400 | 74881 |
| STUDY | 100 | 11400 | 1394 | 8399 |
| GRADE | 651 | 74881 | 8399 | 53617 |

X'X Inverse, Parameter Estimates, and SSE

| Variable | Intercept | IQ | STUDY | GRADE |
|-----------|--------------|--------------|--------------|--------------|
| Intercept | 28.898526711 | -0.226082693 | -0.224182192 | 0.7365546771 |
| IQ | -0.226082693 | 0.0018460178 | 0.0011217122 | 0.473083715 |
| STUDY | -0.224182192 | 0.0011217122 | 0.0076260404 | 2.1034362851 |
| GRADE | 0.7365546771 | 0.473083715 | 2.1034362851 | 45.759884688 |

Analysis of Variance

| Source | DF | Sum of Squares | Mean Square | F Value | Pr > F |
|-----------------|----|-------------------|----------------|---------|--------|
| Model | 2 | 596.11512 | 298.05756 | 32.57 | 0.0014 |
| Error | 5 | 45.75988 | 9.15198 | | |
| Corrected Total | 7 | 641.87500 | | | |

(Output continued next page)

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|--------------------|----------------|---------|---------|
| Intercept | 1 | 0.73655 | 16.26280 | 0.05 | 0.9656 |
| IQ | 1 | 0.47308 | 0.12998 | 3.64 | 0.0149 |
| STUDY | 1 | 2.10344 | 0.26418 | 7.96 | 0.0005 |

Covariance of Estimates

| Variable | Intercept | IQ | STUDY |
|-----------|--------------|--------------|--------------|
| Intercept | 264.47864999 | -2.069103589 | -2.051710248 |
| IQ | -2.069103589 | 0.016894712 | 0.010265884 |
| STUDY | -2.051710248 | 0.010265884 | 0.0697933458 |

Output Statistics

| Obs | Dependent Variable | Predicted Value | Std Error Mean Predict | 95% CL Mean | 95% CL Predict | Residual |
|---------------|--------------------|-----------------|------------------------|-------------|----------------|----------|
| 1 | 75 | 71.4447 | 1.9325 | 66.4770 | 76.4124 | 3.5553 |
| (abbreviated) | | | | | | |
| 8 | 76 | 80.5426 | 1.9287 | 75.5847 | 85.5005 | -4.5426 |
| 9 | . | 83.6431 | 1.1414 | 80.7092 | 86.5771 | . |

| | |
|-------------------------------|-----------|
| Sum of Residuals | 0 |
| Sum of Squared Residuals | 45.75988 |
| Predicted Residual SS (PRESS) | 125.73575 |

(This 9th “observation” in the output data set is an illustration of the “missing y” trick to get software to generate prediction limits.)

```

> # lm() in R
> iqstudy.dat
  IQ study grade
1 105   10    75
2 110   12    79
(abbreviated)
7 114   20    98
8 102   15    76
> iqstudy.out <- lm(iqstudy.dat$grade ~ iqstudy.dat$IQ + iqstudy.dat$study)
> coef(iqstudy.out)
      (Intercept)      iqstudy.dat$IQ iqstudy.dat$study
      0.7365547      0.4730837      2.1034363
> vcov(iqstudy.out)
      (Intercept) iqstudy.dat$IQ iqstudy.dat$study
(Intercept)      264.478650      -2.06910359      -2.05171025
iqstudy.dat$IQ      -2.069104      0.01689471      0.01026588
iqstudy.dat$study  -2.051710      0.01026588      0.06979335
> summary(iqstudy.out)
Call: lm(formula = iqstudy.dat$grade ~ iqstudy.dat$IQ + iqstudy.dat$study)

Residuals:
    1     2     3     4     5     6     7     8 
3.55529  0.98300 -2.12722  2.04106 -1.10775 -0.06493  1.26318 -4.54264 

Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept)    0.7366    16.2628   0.045 0.965629
iqstudy.dat$IQ  0.4731     0.1300   3.640 0.014909 *
iqstudy.dat$study 2.1034     0.2642   7.962 0.000504 ***

Residual standard error: 3.025 on 5 degrees of freedom
Multiple R-squared:  0.9287,    Adjusted R-squared:  0.9002 
F-statistic: 32.57 on 2 and 5 DF,  p-value: 0.001357

```


$$\textcircled{0} \mu(x_1, x_2, x_3) = E(Y|x_1, x_2, x_3) = \beta_0 \quad \text{Variable Selection}$$

x_1, x_2, x_3 denote p independent variables. Consider several models:

$$\textcircled{1} \mu(x_1, x_2, x_3) = E(Y|x_1, x_2, x_3) = \beta_0 + \beta_1 x_1$$

$$\textcircled{2} \mu(x_1, x_2, x_3) = E(Y|x_1, x_2, x_3) = \beta_0 + \beta_2 x_2 \quad \text{SLR of } Y \text{ on } x_2$$

$$\textcircled{3} \mu(x_1, x_2, x_3) = E(Y|x_1, x_2, x_3) = \beta_0 + \beta_3 x_3$$

$$\textcircled{4} \mu(x_1, x_2, x_3) = E(Y|x_1, x_2, x_3) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_3 \quad \text{MLR on } x_1, x_2, x_3 \quad \text{"additive"}$$

$$\textcircled{5} \mu(x_1, x_2, x_3) = E(Y|x_1, x_2, x_3) = \beta_0 + \beta_1 x_1 + \beta_3 x_3$$

$$\textcircled{6} \mu(x_1, x_2, x_3) = E(Y|x_1, x_2, x_3) = \beta_0 + \beta_1 x_1 + \beta_2 x_2$$

$$\textcircled{7} \mu(x_1, x_2, x_3) = E(Y|x_1, x_2, x_3) = \beta_0 + \beta_2 x_2 + \beta_3 x_3$$

Language

A is nested in B means model A can be obtained by placing linear restrictions on parameter values in model B (e.g. $\beta_1 = \beta_2$)

True or false:

- Model 1 nested in Model 4 $\text{True: } (\beta_2 = \beta_3 = 0)$
- Model 2 nested in Model 4 True
- Model 3 nested in Model 4 True

- Model 1 nested in Model 5 $\text{True } (\beta_3 = 0)$
- Model 4 nested in Model 1 False
- Model 5 nested in Model 4 $\text{True } (\beta_2 = 0)$

A nested in B \rightarrow A called *reduced*, B called *full*.

p - number of regression parameters in full model

q - number of regression parameters in reduced model

$p - q$ - number of regression parameters being tested.

$$SS[Total] = SS[R] + SS[E]$$

$$\sum (Y_i - \bar{Y})^2 = \sum (\hat{Y}_i - \bar{Y})^2 + \sum (\hat{Y}_i - Y_i)^2$$

Variable/model Selection - concepts

In comparing two models, suppose

β_1, \dots, β_q in reduced (r) model (A)

$\beta_1, \dots, \beta_q, \beta_{q+1}, \dots, \beta_p$ in full (f) model (B).

Comparison of models A and B amounts to testing

$H_0: \beta_{q+1} = \beta_{q+2} = \dots = \beta_p = 0$ (model A ok) $df = p - q$

$H_1: \beta_{q+1}, \beta_{q+2}, \dots, \beta_p$ not all 0 (need model B)

$$\text{Let } F = \frac{(SS[E]_r - SS[E]_f) / (p - q)}{MS[E]_f} = \frac{MS[H_0]}{MS[E]}$$

Difference in the numerator called an *extra regression sum of squares*:

$$R(\beta_{q+1}, \beta_{q+2}, \dots, \beta_p | \beta_0, \beta_1, \beta_2, \dots, \beta_q) = SS[R]_f - SS[R]_r = \underline{\hspace{2cm}}$$

(ok to suppress β_0 in these extra SS terms.) $= SS(E)_r - SS(E)_f$

Theory gives that if H_0 holds (model A appropriate), F behaves according to F distribution with $p - q$ numerator, $n - p - 1$ denominator degrees of freedom. (Write $\sim F_{p-q, n-p-1}$)

Extra SS terms for comparing some nested models on preceding page:

- Model 1 in model 4: $R(\beta_2, \beta_3 | \beta_1)$
- Model 2 in model 4 ? $R(\beta_1, \beta_3 | \beta_0, \beta_2)$ (H0: $\beta_1 = \beta_3 = 0$)
- Model 3 in model 4 ? $R(\beta_1, \beta_2 | \beta_0, \beta_3)$
- Model 1 in model 5: $R(\beta_3 | \beta_1)$
- Model 5 in model 4: $R(\beta_2 | \beta_0, \beta_1, \beta_3)$

To compare Models 1 and 4, compute $F = (R(\beta_2, \beta_3 | \beta_1)/2)/MSE_4$ on $df = 2, n - 3 - 1$ If observed F sufficiently large, models said to differ significantly, reduced model rejected in favor of full model. If F small, reduced model plausible.

An example: How to measure body fat?

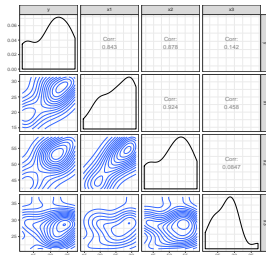
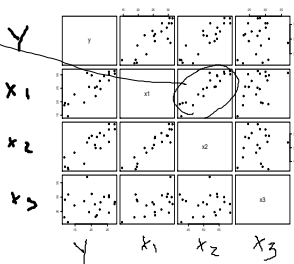
For each of $n = 20$ healthy individuals, the following measurements were made: bodyfat percentage y_i , triceps skinfold thickness, x_1 , thigh circumference x_2 , midarm circumference x_3 . (See "bodyfat.txt")

```
x1    x2    x3    y
19.5  43.1  29.1  11.9
24.7  49.8  28.2  22.8
(abbreviated)
22.7  48.2  27.1  14.8
25.2  51.0  27.5  21.1
```

Summary statistics:

| Symbol | Variable | ^{avg} mean | st. dev. |
|--------|--------------|-----------------------------------|----------|
| y | Body fat | 20.2 | 5.1 |
| x1 | Triceps | 25.3 | 5.0 |
| x2 | Thigh Circ. | 51.2 | 5.2 |
| x3 | Midarm Circ. | 27.6 | 3.6 |

Collinearity



```
> pairs(bodyfat.df[,c(4,1:3)])
> scatterplot(bodyfat.df, lower=list(continuous="density"), data.var=c(4,1:3), diag=list(continuous=
+ "densityDiag"))
```

Pearson Correlation Coefficients, N = 20
Prob > |r| under H0: Rho=0

| | y | x1 | x2 | x3 |
|----|-------------------|-------------------|-------------------|--------------------------|
| y | 1.00000 | 0.84327 <.0001 | 0.87809 <.0001 | 0.14244 <u>0.5491</u> |
| x1 | 0.84327 <.0001 | 1.00000 | 0.92384 <.0001 | 0.45778 0.0424 |
| x2 | 0.87809 <.0001 | 0.92384 <.0001 | 1.00000 | 0.08467 0.7227 |
| x3 | 0.14244 0.5491 | 0.45778 0.0424 | 0.08467 0.7227 | 1.00000 |

$$H_0: \beta_3 = 0$$

$$t = \frac{\hat{\beta}_3}{SE(\hat{\beta}_3)} = \frac{\hat{\beta}_3}{\sqrt{\frac{r^2(n-2)}{(1-r^2)}}}$$

↑
exercise

— (avg over other variables)

Marginal associations between y and x_1 and between y and x_2 are highly significant, providing evidence of a strong $r \approx 0.85$ linear association between bodyfat and triceps skinfold and between bodyfat and thigh circumference.

Multicollinearity: linear associations among the independent variables; causes problems such as inflated sampling variances for $\hat{\beta}$.

x_1 and x_2 are particularly problematic. Imagine trying to balance a planar table top in the third dimension over “legs” that arise from the (x_1, x_2) coordinates. Highly unstable.

```

data bodyfat;
  input x1 x2 x3 y;          cards;
  19.5 43.1 29.1 11.9
  24.7 49.8 28.2 22.8
  (data abbreviated)
  22.7 48.2 27.1 14.8
  25.2 51.0 27.5 21.1
;
proc reg data=bodyfat;
  model y=x1 x2 x3; model y=x1; model y=x2; model y=x3; *usually use 4 lines of code;

```

M4

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|--------------------|----------------|---------|---------|
| Intercept | 1 | 117.08469 | 99.78240 | 1.17 | 0.2578 |
| x1 | 1 | 4.33409 | 3.01551 | 1.44 | 0.1699 |
| x2 | 1 | -2.85685 | 2.58202 | -1.11 | 0.2849 |
| x3 | 1 | -2.18606 | 1.59550 | -1.37 | 0.1896 |

$$\mu(x_1, x_2, x_3) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_3$$

$$\mu(\quad) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_3$$

$$H_0: \beta_1, \beta_2, \beta_3 = 0 \quad H_0: \beta_1 = 0$$

M1

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|--------------------|----------------|---------|---------|
| Intercept | 1 | -1.49610 | 3.31923 | -0.45 | 0.6576 |
| x1 | 1 | 0.85719 | 0.12878 | 6.66 | <.0001 |

$$\mu(x_1) = \beta_0$$

$$\mu(x_1) = \beta_0 + \beta_1 x_1$$

$$H_0: \beta_1 = 0$$

M2

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|--------------------|----------------|---------|---------|
| Intercept | 1 | -23.63449 | 5.65741 | -4.18 | 0.0006 |
| x2 | 1 | 0.85655 | 0.11002 | 7.79 | <.0001 |

M3

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|--------------------|----------------|---------|---------|
| Intercept | 1 | 14.68678 | 9.09593 | 1.61 | 0.1238 |
| x3 | 1 | 0.19943 | 0.32663 | 0.61 | 0.5491 |

Model Selection - examples

In bodyfat data, consider comparing SLR of Y on x_1 with full additive model.

$$\text{Model A : } \mu(x_1, x_2, x_3) = \beta_0 + \beta_1 x_1$$

$$\text{Model B : } \mu(x_1, x_2, x_3) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_3$$

$$\begin{aligned} q &= 1 \\ p &\approx 3 \\ p - q &= 2 \end{aligned}$$

or the null hypothesis

$$\sqrt{H_0 : \beta_2 = \beta_3 = 0 \text{ vs } H_1 : \beta_2, \beta_3 \text{ not both } 0}$$

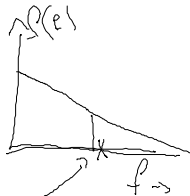
after accounting for x_1 .

$$F = \frac{SS(R)_B - SS(R)_A}{6.15} = \frac{22.3}{6.15} = 3.64$$

How many df ? The 95th percentile is $F(0.05, 2, 46) = 3.63$.

Q: Conclusion from this comparison of nested models?

critical
value



After accounting for effect of x_1 , the (partial) association between y and the pair x_2 and/or x_3 may be declared significant at $\alpha = 0.05$.

```
* To get this F$-ratio in SAS, try ;
proc reg data=bodyfat;
  model y=x1 x2 x3;
  test x2=0, x3=0;
run;
```



```
proc reg;
  model y=x1 x2 x3 / ss1 ss2;
run;
```

The SAS System
The REG Procedure

Analysis of Variance

| Source | DF | Sum of Squares | Mean Square | F Value | Pr > F |
|-----------------|----|----------------|-------------|---------|--------|
| Model | 3 | 396.98461 | 132.32820 | 21.52 | <.0001 |
| Error | 16 | 98.40489 | 6.15031 | | |
| Corrected Total | 19 | 495.38950 | | | |

| | | | |
|----------------|----------|----------|--------|
| Root MSE | 2.47998 | R-Square | 0.8014 |
| Dependent Mean | 20.19500 | Adj R-Sq | 0.7641 |
| Coeff Var | 12.28017 | | |

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t | Type I SS | Type II SS |
|-----------|----|--------------------|----------------|---------|---------|------------|------------|
| Intercept | 1 | 117.08469 | 99.78240 | 1.17 | 0.2578 | 8156.76050 | 8.46816 |
| x1 | 1 | 4.33409 | 3.01551 | 1.44 | 0.1699 | 352.26980 | 12.70489 |
| x2 | 1 | -2.85685 | 2.58202 | -1.11 | 0.2849 | 33.16891 | 7.52928 |
| x3 | 1 | -2.18606 | 1.59550 | -1.37 | 0.1896 | 11.54590 | 11.54590 |

Type I - sequential. Type II - partial p-values are *partial*
 Note agreement between p -values from Type II F tests and p -values from t tests from parameter estimates from MLR.

Type I sums of squares - sequential (order of selection matters)

Type II sums of squares - partial (Δ SSE due to adding term A to model with all other terms not 'containing' A)

Type III sums of squares - partial

extra SS
reg \rightarrow

| | | | |
|--|---|-------|--|
| $R(\beta_1 \beta_0)$ | = | 352.3 | |
| $R(\beta_2 \beta_0, \beta_1)$ | = | 33.17 | |
| $R(\beta_3 \beta_0, \beta_1, \beta_2)$ | = | 11.5 | |
| $R(\beta_1 \beta_0, \beta_2, \beta_3)$ | = | 12.7 | |
| $R(\beta_2 \beta_0, \beta_1, \beta_3)$ | = | 7.5 | |

$MSE_1 =$
 $MSE_2 =$
 $MSE_4 = 6.15$

Type II test for β_j - test of partial association between y and x_j after accounting for all other x_i

Type II F -ratios from bodyfat data for x_1, x_2, x_3 , respectively:

$$F = \frac{12.7/1}{6.15} = 2.07, \quad F = \frac{7.5/1}{6.15} = 1.22, \quad F = \frac{11.5/1}{6.15} = 1.88.$$

(Partial) effects significant ? (Use $F(0.95, 1, 16) = 4.49$.)

Exercise: Carry out the corresponding F -tests to compare models.

In PROC REG output, which models are the type I tests comparing?

- 1 Type I SS for x_1 appropriate for SLR of y on x_1 .
- 2 Type I SS for x_2 appropriate for test of association between y and x_2 after accounting for x_1 . $R(\beta_2 | \beta_0, \beta_1) = 33.2$
- 3 Type I test for x_3 same as type II test for x_3 .

In all three of these tests, $MS[E]$ computed from full model (#4).

Some model comparison examples

- 1 Compare models 1 and 6
- 2 Compare models 2 and 6

For 1. use $R(\beta_2 | \beta_0, \beta_1)$ in the F ratio:

$$\begin{aligned}
 F &= \frac{R(\beta_2 | \beta_0, \beta_1)}{MS[E]_6} \\
 &= \frac{33.2}{(SS[Total] - R(\beta_1, \beta_2 | \beta_0) / (20 - 2 - 1))} \\
 &= \frac{33.2}{(495.4 - 352.3 - 33.2) / (20 - 2 - 1)} \\
 &= \frac{33.2}{109.9 / 17} = 5.1
 \end{aligned}$$

Note that $SS[E]_f = (SS[Total] - SS[R]_f)$ and $SS[R]_f = SS[R]_r + R(\beta_2 | \beta_0, \beta_1)$

$F(0.05, 1, 17) = 4.45$: model 1 rejected in favor of model 6: there is evidence ($p = 0.037$) of association between y and x_2 after accounting for dependence on x_1 .

| Source | df | SS | MS |
|--------|----|-------|----|
| Reg | 1 | 352.3 | |
| Error | 18 | 143.1 | |
| Tot | 19 | 495.4 | |

| Source | df | SS | MS |
|--------|----|-------|------|
| Reg | 2 | 385.5 | |
| Error | 17 | 109.9 | 6.46 |
| Tot | 19 | 495.4 | |

$H_0: \beta_2 = 0$

| M2 ANOVA | | | |
|----------|----|-------|----|
| Source | df | SS | MS |
| x_2 | 1 | 38.2 | |
| Error | 18 | 113.4 | |
| Tot | 19 | 495.4 | |

To compare models 2 and 6, we need $SS[R]_r = R(\beta_2|\beta_0) = 382.0$ which cannot be gleaned from preceding output. You could also get it from $r_{yx_2}^2 \times SS[Tot]$ or from running something like

```
proc reg;
  model y=x1 x2/ss1 ss2;
run;
```

The REG Procedure

| Source | DF | Sum of Squares | Mean Square | F Value | Pr > F |
|-----------------|----|----------------|-------------|---------|--------|
| Model | 2 | 385.43871 | 192.71935 | 29.80 | <.0001 |
| Error | 17 | 109.95079 | 6.46769 | | |
| Corrected Total | 19 | 495.38950 | | | |

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t | Type I SS | Type II SS |
|-----------|----|--------------------|----------------|---------|---------|------------|------------|
| Intercept | 1 | -19.17425 | 8.36064 | -2.29 | 0.0348 | 8156.76050 | 34.01785 |
| x1 | 1 | 0.22235 | 0.30344 | 0.73 | 0.4737 | 352.26980 | 3.47289 |
| x2 | 1 | 0.65942 | 0.29119 | 2.26 | 0.0369 | 33.16891 | 33.16891 |

$$M_2: \mu(x_1, x_2) = \beta_0 + \beta_1 x_1 + \beta_2 x_2$$

testing for ass'n b/w

$E(y|x)$ and x_1
after controlling for x_2

$$H_0: \beta_2 = 0$$

$$\begin{aligned}
 F &= \frac{R(\beta_1|\beta_0, \beta_2)/(\Delta df)}{MS[E]_f} \quad \text{Ho } F_{1,17} \\
 &= \frac{(SS[R]_f - SS[R]_r)/1}{6.5} \\
 &= \frac{352.3 + 33.2 - 382.0}{6.5} = \frac{3.5}{6.5} \approx 0.5
 \end{aligned}$$

No evidence against $H_0: \beta_2 = 0$

Conclusions?

- x_2 gives you a little when you add it to model with x_1
- x_1 gives you nothing when you add it to model with x_2
- Take model with x_2 . (Has higher r^2 too.)
- these comparisons of nested models easy to carry out using TEST statement in PROC REG.

$\mu(I, T) = \beta_0$ Another example, revisiting test scores and study times
Consider this sequence of analyses:

- 1 Regress GRADE on IQ.
- 2 Regress GRADE on IQ and TIME.
- 3 Regress GRADE on TIME IQ TI where $TI = \text{TIME} \times \text{IQ}$.

$H_0: \beta_I = 0$
 $H_a: \beta_I \neq 0$

ANOVA (Grade on IQ)

| SOURCE | DF | SS | MS | F | p-value |
|--------|----|---------|---------|-------|---------|
| IQ | 1 | 15.9393 | 15.9393 | 0.153 | 0.71 |
| Error | 6 | 625.935 | 104.32 | | |

No evidence that IQ has anything to do with grade, but we did not look at study time.
Looking at the *multiple* regression we get

The REG Procedure

Analysis of Variance

| Source | DF | Sum of Squares | Mean Square | F Value | Pr > F |
|-----------------|----|----------------|-------------|---------|--------|
| Model | 2 | 596.11512 | 298.05756 | 32.57 | 0.0014 |
| Error | 5 | 45.75988 | 9.15198 | | |
| Corrected Total | 7 | 641.87500 | | | |

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|--------------------|----------------|---------|---------|
| Intercept | 1 | 0.73655 | 16.26280 | 0.05 | 0.9656 |
| IQ | 1 | 0.47308 | 0.12998 | 3.64 | 0.0149 |
| study | 1 | 2.10344 | 0.26418 | 7.96 | 0.0005 |

Now the test for dependence on IQ is significant $p = 0.0149$. Why?

The interaction model

The REG Procedure
Analysis of Variance

| Source | DF | Sum of Squares | Mean Square | F Value | Pr > F |
|-----------------|----|----------------|-------------|---------|--------|
| Model | 3 | 610.81033 | 203.60344 | 26.22 | 0.0043 |
| Error | 4 | 31.06467 | 7.76617 | | |
| Corrected Total | 7 | 641.87500 | | | |

Parameter Estimates

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t | Type I SS | Type II SS |
|-----------|----|--------------------|----------------|---------|---------|-----------|------------|
| Intercept | 1 | 72.20608 | 54.07278 | 1.34 | 0.2527 | 52975 | 13.84832 |
| IQ | 1 | -0.13117 | 0.45530 | -0.29 | 0.7876 | 15.93930 | 0.64459 |
| study | 1 | -4.11107 | 4.52430 | -0.91 | 0.4149 | 580.17582 | 6.41230 |
| IQ_study | 1 | 0.05307 | 0.03858 | 1.38 | 0.2410 | 14.69521 | 14.69521 |

E

(1.38)²

Model discussion. We call the product $I*S = IQ*STUDY$ an “interaction” term.

$$\hat{G} = 72.21 - 0.13 * I - 4.11 * S + 0.0531(I * S)$$

Now if $IQ = 100$ we get

$$\hat{G} = (72.21 - 13.1) + (-4.11 + 5.31)S$$

and if $IQ = 120$ we get

$$\hat{G} = (72.21 - 15.7) + (-4.11 + 6.37)S.$$

With interaction model, one extra hour of study increases expected grade by 1.20 points for someone with $IQ = 100$ and by 2.26 points for someone with $IQ = 120$. Since interaction not significant, we might go back to simpler “additive” model. (example taken from Dickey’s ST512 notes.)

Some questions about design matrices

Recall three models under consideration for the bodyfat data

$$M_1 : \mu(x_1, x_2, x_3) = \beta_0 + \beta_1 x_1$$

$$M_2 : \mu(x_1, x_2, x_3) = \beta_0 + \beta_2 x_2$$

$$M_6 : \mu(x_1, x_2, x_3) = \beta_0 + \beta_1 x_1 + \beta_2 x_2$$

Q: $MS[E]_{M_6} < MS[E]_{M_1}$ and $MS[E]_{M_6} < MS[E]_{M_2}$ but the partial slopes have larger standard errors in M_6 . Why? _____

Design matrices

$$X_{M_6} = \begin{pmatrix} 1 & 19.5 & 43.1 \\ 1 & 24.7 & 49.8 \\ \vdots & \vdots & \vdots \\ 1 & 25.2 & 51.0 \end{pmatrix} \quad X_{M_1} = \begin{pmatrix} 1 & 19.5 \\ 1 & 24.7 \\ \vdots & \vdots \\ 1 & 25.2 \end{pmatrix}$$

$$(X'X)_{M_6} = \begin{pmatrix} ? & 506.1 & 1023.4 \\ & 13386.3 & 26358.7 \\ & & 52888.0 \end{pmatrix}$$

$$(X'X)_{M_1} = \begin{pmatrix} ? & ? \\ & ? \end{pmatrix} \quad (X'X)_{M_1}^{-1} = \begin{pmatrix} 1.39 & -0.053 \\ & 0.002 \end{pmatrix}$$

$$(X'X)_{M_2} = \begin{pmatrix} ? & ? \\ & ? \end{pmatrix} \quad (X'X)_{M_2}^{-1} = \begin{pmatrix} 5.08 & -0.098 \\ & 0.0019 \end{pmatrix}$$

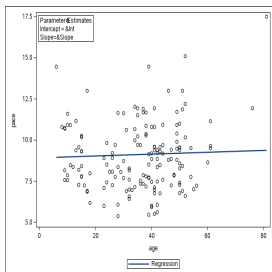
$$(X'X)_{M_6}^{-1} = \begin{pmatrix} 10.8 & 0.29 & -0.35 \\ & 0.014 & -0.012 \\ & & 0.013 \end{pmatrix}$$

Q: Why is $\text{Var}(\hat{\beta}_0)$ bigger in M_2 than in M_1 ?

```

pace    age    sex
5.38333  28     M
5.46667  39     M
(abbreviated)
17.2667  10     F
17.5000  81     M

```



Resolution 5k run, Centennial campus

| Symbol | Variable | mean | st. dev. | variance |
|--------|----------|------|----------|----------|
| y | Pace | 9.1 | 2.2 | 5.0 |
| x | Age | 35.1 | 14.7 | 216.5 |

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|--------------------|----------------|---------|---------|
| Intercept | 1 | 8.92271 | 0.45724 | 19.51 | <.0001 |
| age | 1 | 0.00564 | 0.01203 | 0.47 | 0.6396 |

Let $E_i \stackrel{iid}{\sim} N(0, \sigma^2)$. Quadratic model for pace (Y) as a function of age (x):

$$Y_i = \beta_0 + \beta_1 x_i + \beta_2 x_i^2 + E_i \quad \text{for } i = 1, \dots, 160$$

- $\beta = (\beta_0, \beta_1, \beta_2)'$ is a vector of unknown regression parameters
- σ^2 is the unknown error variance of paces given age x .

Compare this model with the (previously discarded) SLR model

$$Y_i = \beta_0 + \beta_1 x_i + E_i \quad \text{for } i = 1, \dots, 160$$

Q1: Does β_1 have the same interpretation in both models?

Q2: How can we compare the two models?

A2: Using F -ratios to compare nested models (see output next page).

$$\begin{aligned} F &= \frac{R(\beta_2 | \beta_0, \beta_1)}{MS[E]_{full}} \\ &= \frac{(SS[R]_{full} - SS[R]_{red})/1}{MS[E]_{full}} = \frac{(SS[E]_{red} - SS[E]_{full})/1}{MS[E]_{full}} \\ &= \frac{(113.6 - 1.1)/1}{4.3} = \frac{(787.0 - 674.4)/1}{4.3} \\ &= 26.2 \\ &= \left(\frac{\hat{\beta}_2}{SE} \right)^2 \end{aligned}$$

with $F(0.05, 1, 157) = 3.90$. Since $26.2 \gg 3.9$, the linear model is implausible when compared to the quadratic model. Also, $R(\beta_1, \beta_2 | \beta_0) = 113.6$, $F = (113.6/2)/4.3 = 26.4/2 = 13.2$ so that $H_0 : \beta_1 = \beta_2 = 0$ can be rejected.

```

PROC REG DATA=one; /* age2 defined in data step as age*age */
  MODEL pace=age; /* not necessary in light of MODEL2 statement */
  MODEL pace=age age2/ss1; /* ss1 generates sequential sums of squares */
RUN;

```

The REG Procedure
Model: MODEL1

Analysis of Variance

| Source | DF | Sum of Squares | Mean Square | F Value | Pr > F |
|-----------------|-----|-------------------|----------------|---------|--------|
| Model | 1 | 1.09650 | 1.09650 | 0.22 | 0.6396 |
| Error | 158 | 786.99821 | 4.98100 | | |
| Corrected Total | 159 | 788.09472 | | | |

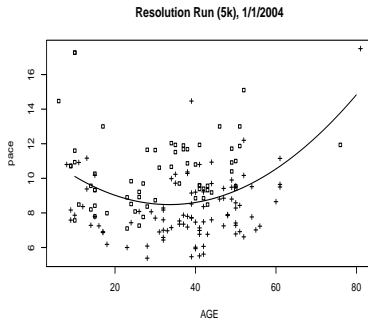
| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|-----------------------|-------------------|---------|---------|
| Intercept | 1 | 8.92271 | 0.45724 | 19.51 | <.0001 |
| age | 1 | 0.00564 | 0.01203 | 0.47 | 0.6396 |

Model: MODEL2

Analysis of Variance

| Source | DF | Sum of Squares | Mean Square | F Value | Pr > F |
|-----------------|-----|-------------------|----------------|---------|--------|
| Model | 2 | 113.64500 | 56.82250 | 13.23 | <.0001 |
| Error | 157 | 674.44972 | 4.29586 | | |
| Corrected Total | 159 | 788.09472 | | | |

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t | Type I SS |
|-----------|----|-----------------------|-------------------|---------|---------|-----------|
| Intercept | 1 | 11.78503 | 0.70216 | 16.78 | <.0001 | 13310 |
| age | 1 | -0.19699 | 0.04113 | -4.79 | <.0001 | 1.09650 |
| age2 | 1 | 0.00294 | 0.00057380 | 5.12 | <.0001 | 112.54850 |



Fitted model is

$$\hat{\mu}(x) = 11.785 - 0.197 x + 0.00294 x^2$$

or

$$\hat{\mu}(\text{age}) = 11.785 - 0.197 \text{ age} + 0.00294 \text{ age}^2.$$

Inference for response Y given predictor x_i .

Random sample of $n = 31$ trees drawn from population of trees. $p = 3$ variables measured on each:

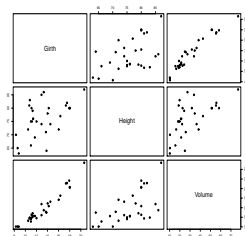
- x_{i1} : "girth", tree diameter in inches
- x_{i2} : "height" (in feet)
- Y_i : volume of timber, in cubic feet.

Given x_1 and x_2 , a MLR model for these data given by

$$Y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + E_i \text{ for } i = 1, \dots, n$$

For trees with x_1, x_2 the model for mean volume is

$$\mu(x_1, x_2) = E(Y|x_1, x_2) = \beta_0 + \beta_1 x_1 + \beta_2 x_2.$$

A scatterplot matrix

Some questions involving linear combinations of regression coefficients

Consider all trees with girth $x_{01} = 15$ in and height $x_{02} = 80$ ft .

- Estimate the mean volume among these trees, along with a standard error and 95% confidence interval.
- Obtain a 95% prediction interval of y_0 , the volume from an individual tree sampled from this population of 80 footers, with girth 15 inches.

SAS generates $\hat{\beta}$ and $\widehat{Var}(\hat{\beta}) = MSE * (X'X)^{-1}$

| Source | DF | Sum of Squares | Mean Square | F Value | Pr > F |
|-----------------|----|----------------|-------------|---------|--------|
| Model | 2 | 7684.16251 | 3842.08126 | 254.97 | <.0001 |
| Error | 28 | 421.92136 | 15.06862 | | |
| Corrected Total | 30 | 8106.08387 | | | |

| | | | |
|----------------|----------|----------|--------|
| Root MSE | 3.88183 | R-Square | 0.9480 |
| Dependent Mean | 30.17097 | Adj R-Sq | 0.9442 |
| Coeff Var | 12.86612 | | |

Parameter Estimates

| Variable | DF | Parameter Estimate | Standard Error | t Value | Pr > t |
|-----------|----|--------------------|----------------|---------|---------|
| Intercept | 1 | -57.98766 | 8.63823 | -6.71 | <.0001 |
| Girth | 1 | 4.70816 | 0.26426 | 17.82 | <.0001 |
| Height | 1 | 0.33925 | 0.13015 | 2.61 | 0.0145 |

Covariance of Estimates

| Variable | Intercept | Girth | Height |
|-----------|--------------|--------------|--------------|
| Intercept | 74.6189461 | 0.4321713812 | -1.050768886 |
| Girth | 0.4321713812 | 0.0698357838 | -0.017860301 |
| Height | -1.050768886 | -0.017860301 | 0.0169393298 |

Inference for the mean response in MLR Recall that if Y a $p \times 1$ random vector with mean μ and covariance matrix Σ . and a a $p \times 1$ (fixed) vector of coefficients.

$$\begin{aligned} E(a'Y) &= a'\mu \\ \text{Var}(a'Y) &= a'\Sigma a. \end{aligned}$$

Consider subpopulation of trees with Girth 15 and Height 80. To estimate mean volume among these trees, with estimated std. error, take $x_0' = (1, 15, 80)$ and consider $\hat{\mu}(x_0) = x_0'\hat{\beta}$.

$$\begin{aligned} E(x_0'\hat{\beta}) &= x_0'\beta \\ \text{Var}(x_0'\hat{\beta}) &= x_0'\hat{\Sigma}x_0 \end{aligned}$$

Substitution of $\hat{\beta}$ and $\hat{\Sigma} = MSE(X'X)^{-1}$ gives the estimates:

$$\begin{aligned} \hat{\mu}(x_0) &= (1, 15, 80) \begin{pmatrix} -58.0 \\ 4.71 \\ 0.34 \end{pmatrix} = 39.8 \\ \widehat{\text{Var}}(\hat{\mu}(x_0)) &= (1, 15, 80) \begin{pmatrix} 74.62 & 0.43 & -1.05 \\ 0.43 & 0.070 & -0.018 \\ -1.05 & -0.018 & 0.017 \end{pmatrix} \begin{pmatrix} 1 \\ 15 \\ 80 \end{pmatrix} = 0.72 \\ \widehat{SE}(\hat{\mu}(x_0)) &= \sqrt{.72} = 0.849 \end{aligned}$$

which can be obtained using PROC REG and the missing y trick:

| Obs | treenumber | Girth | Height | Volume | p | sepred |
|-----|------------|-------|--------|--------|---------|---------|
| 32 | 100 | 15 | 80 | . | 39.7748 | 0.84918 |

95% Prediction limits? Use $\pm t(.025, 28)\sqrt{.72 + MS(E)}$.

Partial correlations

The partial correlation coefficient for x_1 in the MLR

$$E(Y|x_1, x_2, \dots, x_p) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots + \beta_p x_p$$

is defined as the correlation coefficient between the residuals computed from the two regressions below:

$$Y = \beta_0 + \beta_2 x_2 + \dots + \beta_p x_p + E$$

$$X_1 = \beta_0 + \beta_2 x_2 + \dots + \beta_p x_p + E$$

Call these sets of residuals $e_{y \cdot 2, 3, \dots, p}$ and $e_{1 \cdot 2, 3, \dots, p}$ respectively. The *partial correlation* between y and x_1 after accounting for the linear association between y and x_2, x_3, \dots, x_p is defined as

$$r_{y1 \cdot 2, 3, \dots, p} = \text{correlation between } e_{y \cdot 2, 3, \dots, p} \text{ and } e_{1 \cdot 2, 3, \dots, p}.$$

The *partial coeff. of determination* is $r_{y1 \cdot 2, 3, \dots, p}^2$.

Note also that

$$r_{y1 \cdot 2, \dots, p}^2 = \frac{R(\beta_1 | \beta_0, \beta_2, \dots, \beta_p)}{SS[Total] - R(\beta_2, \beta_3, \dots, \beta_p | \beta_0)}.$$

Bodyfat data, compare models 1,2 and 6 (ignore x_3 .)

| bodyfat data | | | | | | | | | |
|---------------|------|------|------|---------|----------|----------|---------|----------|---------|
| Obs | x1 | x2 | y | py_1 | ey_1 | e2_1 | py_2 | ey_2 | e1_2 |
| 1 | 19.5 | 43.1 | 11.9 | 15.2190 | -3.31903 | -2.48145 | 13.2827 | -1.38267 | 1.34939 |
| 2 | 24.7 | 49.8 | 22.8 | 19.6764 | 3.12360 | -0.78756 | 19.0215 | 3.77847 | 0.60956 |
| (abbreviated) | | | | | | | | | |
| 20 | 25.2 | 51.0 | 21.1 | 20.1050 | 0.99500 | -0.06892 | 20.0494 | 1.05061 | 0.04571 |

The partial correlation coefficient between y and x_1 after accounting for x_2 is $r_{y1.2} = 0.17$ and the partial for x_2 after accounting for x_1 is $r_{y2.1} = 0.48$. The partial coefficients of determination are

$$r_{y1.2}^2 = 0.03062 \text{ and } r_{y2.1}^2 = 0.23176.$$

Q: If you had to choose one variable or the other from x_1 and x_2 , which would it be?

Q: Anything wrong with throwing both x_1 and x_2 in the final model?

Q: Write coefficients of determination in terms of extra sums of squares. Use $R(\cdot|\cdot)$ notation.

Note: partial correlations obtained in SAS using PCORR2 option:

| Variable | DF | Squared Partial Corr Type II |
|-----------|----|------------------------------------|
| Intercept | 1 | . |
| x1 | 1 | 0.03062 |
| x2 | 1 | 0.23176 |

Partial regression plots

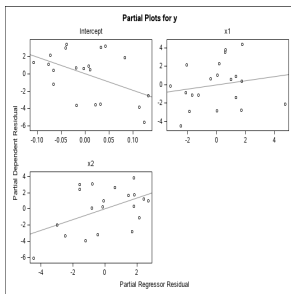
A plot of the residuals from the regression

$$Y = \beta_0 + \beta_2 x_2 + \cdots + \beta_p x_p + E$$

versus the residuals from the regression

$$X_1 = \beta_0 + \beta_2 x_2 + \cdots + \beta_p x_p + E$$

is called a partial regression or leverage plot for x_1 or in the MLR. Can be generated using ODS GRAPHICS ON and the PARTIAL command in the MODEL statement of PROC REG:



Q: What can these plots tell us?

A1: They convey info. about linear associations between y and candidate variable x_i after accounting for linear dependence of y on other variables $x_1, x_2, \dots, x_{i-1}, x_{i+1}, \dots, x_p$

A2: can convey info about nonlinear associations between y and x_i after accounting for other linear associations

A3: They can illuminate possible outliers.

Some exercises (hint: use matrix algebra or SAS).

- ➊ Regarding the `resrun` data as randomly sampled a population of interest. Consider the sub-population of 32 year old males. Fit a quadratic regression function and use it to obtain an estimate of the mean 5k pace in this cohort of all 32 yr-old male runners. Report a standard error and 95% confidence interval.
- ➋ Obtain a 95% prediction interval for one such runner.
- ➌ Explain the difference between the two intervals in questions 1 and 2.
- ➍ At what rate is $\mu(x)$ changing with age? Estimate the appropriate function.
- ➎ Estimate θ , the peak age to run a 5k in the fastest time. Is θ a linear function of regression parameters? Can you obtain an unbiased estimate of the standard error of θ ?

Cook's D

Cook's D for an observation i is a measure of influence on predictions:

$$D_i = \frac{\sum_{j=1}^n (\hat{y}_j - \hat{y}_{j(i)})^2}{pMS(E)}$$

where $\hat{y}_{j(i)}$ is the fitted value for the j^{th} observation if the i^{th} observation is held out.

cdp from ODS output

| Obs | pace | age | sexf | CooksD |
|-----|---------|-----|------|---------|
| 1 | 17.5000 | 81 | 0 | 0.52957 |
| 2 | 11.9333 | 76 | 1 | 0.45148 |
| 3 | 17.2667 | 10 | 1 | 0.13515 |

The MEANS Procedure

Analysis Variable : diff

| Mean | Corrected SS | USS |
|-----------|--------------|-----------|
| 0.0271229 | 6.6410416 | 6.7587459 |

mycooksD computed just for age=81 subject

| Obs | _TYPE_ | _FREQ_ | uss | mycooksD |
|-----|--------|--------|---------|----------|
| 1 | 0 | 160 | 6.75875 | 0.52957 |

```

*resruncooksds.sas;
*Cook's D calculated by PROC REG and from definition;
*inspect code carefully;
data one; set one;
    race=1*scan(crace,1,':')+1/60*scan(crace,2,':');
    pace=1*scan(cpace,1,':')+1/60*scan(cpace,2,':');
    /* The SCAN command extracts characters from a character string */
    age2=age*age; sexf=(sex="F"); agef=age*sexf; age2f=age2*sexf;
    pace2=pace; if age=81 then pace2=.;
run;
ods graphics on; ods trace on; ods listing close;
proc reg data=one plots=cooksds;
    id name pace age sexf;
    model pace=age age2 sexf/influence;
    ods output cooksplot=cdp(rename=(id1=name id2=pace id3=age id4=sexf)) ;
    output out=preds1 p=p1;
run;
proc reg data=one; *where age<81;
    model pace2=age age2 sexf/influence;
    output out=preds2 p=p2;
run;
ods listing ;
proc sort data=cdp; by descending cooksds;
proc print data=cdp (obs=3); title "cdp from ODS output";
    var pace age sexf cooksds;
run;

```

Cook's D for every observation may be “delivered” using PROC REG and SAS ODS with the keyword “cooksplot” discovered using [ods trace on](#).

cdp from ODS output

| Obs | pace | age | sexf | CooksD |
|-----|---------|-----|------|---------|
| 1 | 17.5000 | 81 | 0 | 0.52957 |
| 2 | 11.9333 | 76 | 1 | 0.45148 |
| 3 | 17.2667 | 10 | 1 | 0.13515 |

Calculating Cook's D from the definition ...

```

proc sort data=preds1; by name; run;
proc sort data=preds2; by name; run;
data both;
    merge preds1 preds2;
    by name;
    diff=p1-p2;
run;
proc sort data=both; by descending age;run;
proc means data=both mean css uss;
    var diff;
    output out=uss uss=uss;
run;
data uss; set uss; mycooksD=uss/(4*3.19068); run;
/* MSE=3.19068 was hard-coded after inspection of output from MODEL1*/
proc print data=uss;
    title "mycooksD computed just for age=81 subject";
run;

```

The MEANS Procedure

Analysis Variable : diff

| Mean | Corrected SS | USS |
|-----------|--------------|-----------|
| 0.0271229 | 6.6410416 | 6.7587459 |

mycooksD computed just for age=81 subject

3

| Obs | _TYPE_ | _FREQ_ | uss | mycooksD |
|-----|--------|--------|---------|----------|
| 1 | 0 | 160 | 6.75875 | 0.52957 |

More about model selection: R^2 , R_a^2 and Mallows's C_p

In statistical modelling in general, one goal is often to identify a model explains variability in some response (y) of interest through its association with explanatory factors or variables. The principle of model parsimony dictates that it is best to construct a model which explains things, but with as few variables as possible.

Suppose that the true regression function underlying observed data is given by

$$E(Y|x_1, \dots, x_{q+1}) = \beta_0 + \beta_1 x_1 + \dots + \beta_q x_q \quad (1)$$

but that an analysis leads to the model

$$E(Y|x) = \hat{\beta}_0 + \hat{\beta}_1 x_1 + \dots + \hat{\beta}_{q-2} x_{q-2} \quad (2)$$

Model (2) is said to be *underspecified*. On the other hand, suppose another analysis leads to the model

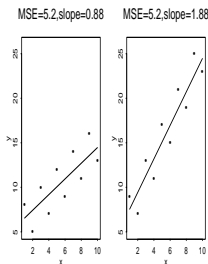
$$E(Y|x) = \hat{\beta}_0 + \hat{\beta}_1 x_2 + \dots + \hat{\beta}_q x_q + \hat{\beta}_{q+1} x_{q+1} \quad (3)$$

Then model (3) is said to be *overspecified*.

- Underspecified models introduce bias ($E(\hat{\mu}(x)) \neq E(Y|x)$.)
- overspecified models inflate the sampling variances of estimators.

Which mistake is worse? (In the sense that underspecification is equivalent to fewer type I errors for tests of the form $H_0 : \beta_i = 0$, it may be preferable, as a rule of thumb.)
The coefficient of multiple determination in MLR, R^2 :

- proportion of variability accounted for by a linear model, also the squared correlation between observed (y_1, y_2, \dots) and predicted $(\hat{y}_1, \hat{y}_2, \dots)$ values
- a reasonable criterion for model selection, but not infallible.



For two datasets with “equal” variability unexplained by SLR, the model with larger absolute slope will have higher R^2 . To see this, recall that

$$R^2 = \frac{SS[R]}{SS[Tot]} = \frac{SS[Tot] - SS[E]}{SS[Tot]} = 1 - \frac{SS[E]}{SS[Tot]}.$$

bigger spread in $y \implies$ bigger R^2

Q: Which line yields a higher r^2 ? Is this a better fit?

R_a^2 , or the *adjusted coefficient of multiple correlation* is given by

$$R_a^2 = 1 - \left(\frac{n-1}{n-p-1} \right) \frac{SS[E]}{SS[Tot]}$$

It imposes a penalty on added independent variables.

Mallow's C_p statistic

Suppose m denotes number of independent variables in full model, $p \leq m$ denotes number of candidates under consideration in reduced model and n denotes sample size.

$$C_p = p + 1 + \frac{(MS[E](p) - MS[E](m)) * (n - p - 1)}{MS[E](m)}$$

Subset models for which $C_p \leq p + 1$ are preferred.

In addition to R^2 , C_p adjusted R^2 , we have AIC, AICc, BIC, ...

Let the likelihood function for a model parameterized by k -dimensional θ (including intercept, if any), using a sample of size n be denoted \mathcal{L}

$$\mathcal{L}(\theta) = f(y_1, \dots, y_n; \theta)$$

Let the maximum likelihood estimator of θ be denoted $\hat{\theta}$. Then

$$AIC = -2 \log \mathcal{L}(\hat{\theta}) + 2k$$

When sample size n is small there is a corrected version of AIC:

$$AICc = AIC + \frac{2k^2 + 2k}{n - k - 1}$$

Schwarz or Bayesian Information Criterion (SIC/BIC)

$$BIC = -2 \log \mathcal{L}(\hat{\theta}) + \log(n)k$$

(penalty in BIC larger than in AIC)

Mallow's C_p for reduced model of dimension q .

$$\begin{aligned} C_p &= \frac{SS[E]_r}{MS[E]_f} + 2p - n \\ &= p + \frac{MS(E)_r - MS(E)_f}{MS(E)_f} (n - p) \end{aligned}$$

Values small $MS(E)_r$ but also small p .

```

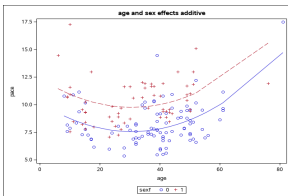
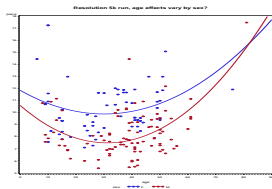
/* models with potentially nonlinear age effects allowed to vary with sex */
proc reg data=one;
  model pace=age age2 sexf agef age2f/selection=cp aic bic rsquare;
run;

```

The SAS System
 The REG Procedure
 Dependent Variable: pace
 C(p) Selection Method

Number of Observations Used

| Number in Model | C(p) | R-Square | 160 Adjusted R-Square | AIC | BIC | MSE | Variables in Model |
|-----------------|---------|----------|-----------------------------|----------|----------|---------|--------------------------|
| 3 | 2.9901 | 0.3684 | 0.3563 | 189.5866 | 191.8431 | 3.19068 | age age2 sexf |
| 4 | 4.8893 | 0.3688 | 0.3525 | 191.4825 | 193.8103 | 3.20918 | age age2 sexf age2f |
| 4 | 4.9883 | 0.3684 | 0.3521 | 191.5848 | 193.9061 | 3.21123 | age age2 sexf agef |
| 4 | 5.0408 | 0.3682 | 0.3519 | 191.6390 | 193.9568 | 3.21232 | age age2 agef age2f |
| 5 | 6.0000 | 0.3725 | 0.3521 | 192.5612 | 195.0257 | 3.21147 | age age2 sexf agef age2f |
| 3 | 11.7363 | 0.3328 | 0.3199 | 198.3700 | 200.1864 | 3.37073 | age age2 agef |
| (abbreviated) | | | | | | | |
| 1 | 89.0585 | 0.0014 | -.0049 | 258.8883 | 259.2590 | 4.98100 | age |



For MLR with i.i.d. normal errors, and $\beta((p+1) \times 1)$

$$\mathcal{L}(\beta, \sigma^2) = \prod_1^n \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left\{-\frac{1}{2}\sigma^2[(y_i - x_i\beta)/\sigma]^2\right\}$$

$$\hat{\beta} = (X'X)^{-1}X'Y$$

$$\hat{\sigma}^2 = \frac{n - (p+1)}{n} MS(E)$$

$$\log \mathcal{L}(\beta, \sigma^2) = -\frac{n}{2} \log 2\pi - \frac{n}{2} \log \sigma^2 - \frac{1}{2\sigma^2} (Y - X\beta)'(Y - X\beta)$$

$$\begin{aligned} \log \mathcal{L}(\hat{\beta}, \hat{\sigma}^2) &= -\frac{n}{2} \log 2\pi - \frac{n}{2} \log \hat{\sigma}^2 - \frac{1}{2\hat{\sigma}^2} (Y - X\hat{\beta})'(Y - X\hat{\beta}) \\ &= -\frac{n}{2} \log \hat{\sigma}^2 + \text{constant} \end{aligned}$$

PROC REG reports $AIC = n \log \left(\frac{SSE}{n} \right) + 2(p+1)$.

[link to SAS PROC REG DOC](#)

For the full model, $MS(E) = 3.21147$ and

```
> 160*(log(3.21147*(160-6)/160)) + 2*6
[1] 192.5612
```

So, check your software's computations!

Residual diagnostics

- Residuals can be plotted against independent/predictor variables to check for model inadequacy. (e.g. if relationship is quadratic, but only a linear model was fit, this plot will reveal a pattern between residuals and predictor.)
- Residuals can be plotted against predicted values to look for inhomogeneity of variance (heteroscedasticity). Look for residuals for which variability increases or “fans out” as one looks left-to-right in this plot (or vice-versa).
- The sorted residuals can be plotted against the normal inverse of the empirical CDF of the residuals in a normal plot to assess the normal distributional assumption. A nonlinear association in such a q-q plot indicates nonnormality. If data-rich, a histogram of residuals can also be used.

Normal plots of residuals

- 1 Obtain the observed quantiles by ordering the residuals:

$$e_{(1)} \leq e_{(2)} \leq \cdots \leq e_{(n)}.$$

- 2 For each $i = 1, \dots, n$ compute the expected quantile from

$$q_{(i)} = z\left(1 - \frac{i}{n+1}\right).$$

- 3 Plot the (ordered) residuals on the vertical axis versus the (ordered) theoretical quantiles on the horizontal axis.

The *empirical cumulative probability* associated with $e_{(i)}$ is

$$p_{(i)} = \frac{\text{Rank of } e_{(i)}}{n+1}.$$

Corresponding theoretical quantiles obtained via

$$q_{(i)} = z(1 - p_{(i)}).$$

e.g. suppose $n = 9$ then for $i = 1$ we look up the 10th percentile of $N(0, 1)$ which is -1.282
 ... for $i = 9$ we look up $q_{(9)} = +1.282$. Plot the ordered residuals (empirical quantiles) against the theoretical quantiles and expect linearity.

```

ods listing close;
ods graphics on;
proc reg data=running;
    model pace=sexf age age2;    *general linear model.  will discuss soon;
    output out=resids p=yhat r=resid;
run;
proc rank data=resids out=resids2;
    ranks rankresid;
    var resid;
run;
data resids2;
    set resids2;
    ecdf=rankresid/(160+1);      *160 runners;
    q=probit(ecdf);
run;
ods listing ;
proc print data=resids2 ;
    var age pace yhat resid rankresid ecdf q;
run;
proc gplot data=resids;
    plot resid*q;
run;

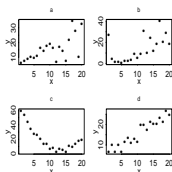
```

The SAS System

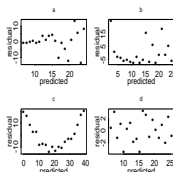
1

| Obs | age | pace | yhat | resid | rankresid | ecdf | q |
|---------------|-----|---------|---------|----------|-----------|---------|----------|
| 1 | 28 | 5.3833 | 7.5837 | -2.20040 | 14.0 | 0.08696 | -1.35974 |
| 2 | 39 | 5.4667 | 7.7671 | -2.30046 | 10.0 | 0.06211 | -1.53728 |
| 3 | 41 | 5.5167 | 7.8735 | -2.35681 | 6.0 | 0.03727 | -1.78332 |
| 4 | 42 | 5.6167 | 7.9351 | -2.31841 | 9.0 | 0.05590 | -1.59015 |
| 5 | 40 | 5.9333 | 7.8175 | -1.88416 | 18.0 | 0.11180 | -1.21700 |
| (abbreviated) | | | | | | | |
| 156 | 6 | 14.4667 | 11.4534 | 3.01324 | 155.0 | 0.96273 | 1.78332 |
| 157 | 52 | 15.1000 | 11.0579 | 4.04215 | 157.0 | 0.97516 | 1.96263 |
| 158 | 10 | 17.2667 | 10.9473 | 6.31937 | 158.5 | 0.98447 | 2.15636 |
| 159 | 10 | 17.2667 | 10.9473 | 6.31937 | 158.5 | 0.98447 | 2.15636 |
| 160 | 81 | 17.5000 | 14.7178 | 2.78223 | 152.0 | 0.94410 | 1.59015 |

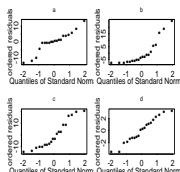
A fun exercise: Match up letters a,b,c,d with the model violation



$y \text{ v } x$



resid v predicted ($y \text{ v } \hat{y}$)



normal plots of resids

- 1 Heteroscedasticity (nonconstant _____)
- 2 Nonlinearity ($\mu(x)$ not linear in _____)
- 3 Nonnormality (vertical variation in y about $\mu(x)$ not _____-shaped)
- 4 Model fits (hurray!)