Mote Regtession using Mattices

Fitted Values

Let the vector of **fitted values** be

$$\hat{\mathbf{Y}}_{n imes 1} = egin{bmatrix} Y_1 \ \hat{Y}_2 \ dots \ \hat{Y}_n \end{bmatrix}$$

In matrix notation

$$\hat{\mathbf{Y}}_{n\times 1} = \mathbf{X}_{n\times 2} \mathbf{b}_{2\times 1}$$

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Substituting the expression for **b**

$$\hat{\mathbf{Y}}_{n\times 1} = \mathbf{X}_{n\times 2} \; (\mathbf{X}'\mathbf{X})^{-1} \; \mathbf{X}' \; \mathbf{Y}_{n\times 1}$$

This is often expressed as

$$\hat{\mathbf{Y}}_{n\times 1} = \mathbf{H}_{n\times n} \mathbf{Y}_{n\times 1}$$

where

$$\mathbf{H}_{n\times n} = \mathbf{X}_{n\times 2} \; (\mathbf{X'X})^{-1} \; \mathbf{X'}_{2\times n}$$

H is a square *n* x *n* matrix that can be created completely from the design matrix and its transpose.

The fitted values can be expressed as linear combinations of the response vector **Y** using coefficients found in **H**.

```
> h %*% Y

     [,1]
1 12.081494
2 12.580862
3 12.081494
4 12.580862
5 9.085287
```

Because of this, **H** is often referred to as the *hat matrix*.

H is a symmetric matrix and also has a special property called *idempotency*:

$$egin{array}{ccc} \mathbf{H} & \mathbf{H} & = & \mathbf{H} \\ n imes n & n imes n \end{array}$$

```
> h %*% h
1 2 3 4 5
1 0.21839080 0.24137931 0.21839080 0.24137931 0.088045977
2 0.24137931 0.29310345 0.24137931 0.29310345 -0.06896552
3 0.21839080 0.24137931 0.21839080 0.24137931 0.08045977
4 0.24137931 0.29310345 0.24137931 0.29310345 -0.06896552
5 0.08045977 -0.06896552 0.08045977 -0.06896552 0.97701149
```

Residuals

Let the vector of residuals be

$$\mathbf{e}_{n \times 1} = \begin{bmatrix} e_1 \\ e_2 \\ \vdots \\ e_n \end{bmatrix}$$

In matrix notation

$$\mathbf{e}_{n\times 1} = \mathbf{Y}_{n\times 1} - \mathbf{X}_{n\times 2} \mathbf{b}_{2\times 1}$$
$$= \mathbf{Y}_{n\times 1} - \mathbf{\hat{Y}}_{n\times 1}$$

Since the fitted values can be expressed as **HY**, the residual vector can also be expressed as

$$\mathbf{e}_{n\times 1} = \mathbf{Y}_{n\times 1} - \mathbf{H}_{n\times n} \mathbf{Y}_{n\times 1}$$

The residuals can also be expressed as linear combinations of the response vector **Y**.

Using the distributive property as it relates to matrices, we re-write this as

$$\mathbf{e}_{n\times 1} = (\mathbf{I}_{n\times n} - \mathbf{H}_{n\times n}) \mathbf{Y}_{n\times 1}$$

The matrix (1 - H), like H, is *symmetric* and *idempotent*.

$$\mathbf{e}_{_{n imes 1}} = (\mathbf{I}_{_{n imes n}} - \mathbf{H}_{_{n imes n}}) \mathbf{Y}_{_{n imes 1}}$$

```
> i = diag(5)
> (i - h) %*% Y
        [,1]
1 -0.08149425
2 -4.58086207
3 4.17850575
4 1.06913793
5 -0.58528736
> resid(lm.1)
-0.08149425 -4.58086207 4.17850575 1.06913793 -0.58528736
```

Variance—Covariance Matrix of the Residuals

Since $\mathbf{e} = (\mathbf{I} - \mathbf{H})\mathbf{Y}$, the variance–covariance matrix of \mathbf{e} is can be expressed as

$$\sigma^2\{\mathbf{e}\} = \sigma^2\{(\mathbf{I}_{n\times n} - \mathbf{H}_{n\times n}) \mathbf{Y}_{n\times 1}\}$$

Using rules of variances, the right-hand side is re-expressed as

$$oldsymbol{\sigma}^2\{\mathbf{e}\} = (\mathbf{I}_{n imes n} - \mathbf{H}_{n imes n}) \ oldsymbol{\sigma}^2\{\mathbf{Y}_{n imes 1}\} \ (\mathbf{I}_{n imes n} - \mathbf{H}_{n imes n})'$$

Now, for the model with normally distributed errors,

$$\sigma^2\{Y\} = \sigma^2\{e\} = \sigma^2 I$$

Also, (I - H) = (I - H)' because of the symmetry of this matrix. Hence,

$$\boldsymbol{\sigma}^2\{\mathbf{e}\} = (\mathbf{I}_{n \times n} - \mathbf{H}_{n \times n}) \ \sigma^2\{\mathbf{I}_{n \times n}\} \ (\mathbf{I}_{n \times n} - \mathbf{H}_{n \times n})$$

Which reduces to

$$\boldsymbol{\sigma}^{2}\{\mathbf{e}\} = \sigma^{2} \left(\mathbf{I}_{n \times n} - \mathbf{H}_{n \times n} \right) \left(\mathbf{I}_{n \times n} - \mathbf{H}_{n \times n} \right)$$

Since (1 - H) is idempotent, (1 - H)(1 - H) = (1 - H) and thus

$$\boldsymbol{\sigma}^2 \{ \mathbf{e} \} = \sigma^2 \left(\mathbf{I}_{n \times n} - \mathbf{H}_{n \times n} \right)$$

$$\sigma^2\{\mathbf{e}\} = \sigma^2 \left(\mathbf{I}_{n \times n} - \mathbf{H}_{n \times n} \right)$$

The variance of residual e_i is

$$\sigma^2\{e_i\} = \sigma^2(1 - h_{ii})$$

where h_{ii} is the i^{th} element on the main diagonal of the hat matrix

The covariance between residuals e_i and e_j ($i \neq j$) is

$$\sigma\{e_i, e_j\} = \sigma^2(0 - h_{ij})$$
$$= -h_{ij}\sigma^2$$

where h_{ij} is the element in the i^{th} row and j^{th} column of the hat matrix

When the MSE is used as an estimator of σ^2

$$s^2\{e_i\} = \text{MSE}(1 - h_{ii})$$

$$s\{e_i, e_j\} = -h_{ij}MSE$$

For our data

$$s^{2}{e_{1}} = 15.33642(1 - 0.33064516) = 10.26551$$

$$s\{e_1, e_2\} = -0.33064516(15.33642) = -5.070913$$

The main diagonal element h_{ii} of the hat matrix can be computed directly as

$$h_{ii} = \mathbf{X}_{i}' \left(\mathbf{X}' \mathbf{X} \right)^{-1} \mathbf{X}_{i}$$

$$_{1 \times p} \sum_{i \in \mathbb{N}} \mathbf{X}_{i} \sum_{n \times p} \mathbf{X}_{i}$$

where X_i is the vector of predictor values for the i^{th} case defined as

$$\mathbf{X}_i = egin{bmatrix} 1 \ X_{i,1} \ dots \ X_{i,p-1} \end{bmatrix}$$

$$h_{ii} = \mathbf{X}_{i}' \left(\mathbf{X}' \mathbf{X} \right)^{-1} \mathbf{X}_{i}$$

$${}_{1 \times p} {}_{1 \times p} {}_{1 \times p} {}_{1 \times p}$$

```
# Obtain the predictor vector for the 1st observation
> X1 = X[1, ]
> h1 = t(X1) %*% solve(t(X) %*% X) %*% X1
> h1

[,1]
[1,] 0.2183908
```

$$s^2\{e_i\} = MSE(1 - h_{ii})$$

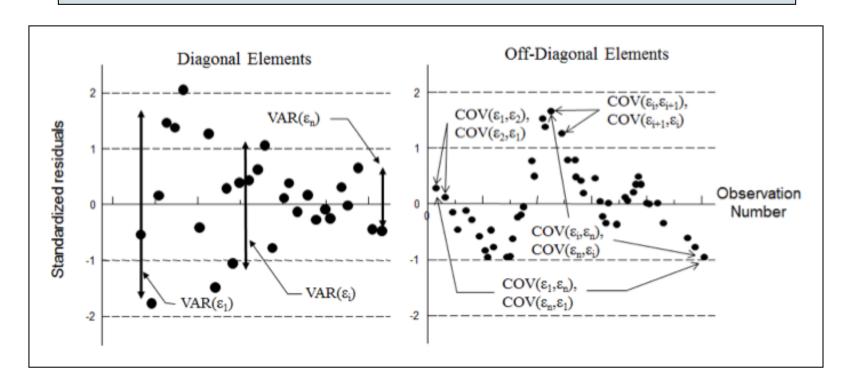
```
# Compute the residual variance estimate for the 1st observation
> MSE * (1 - h1)

[,1]
[1,] 10.40478
```

Recall the estimated variance—covariance matrix

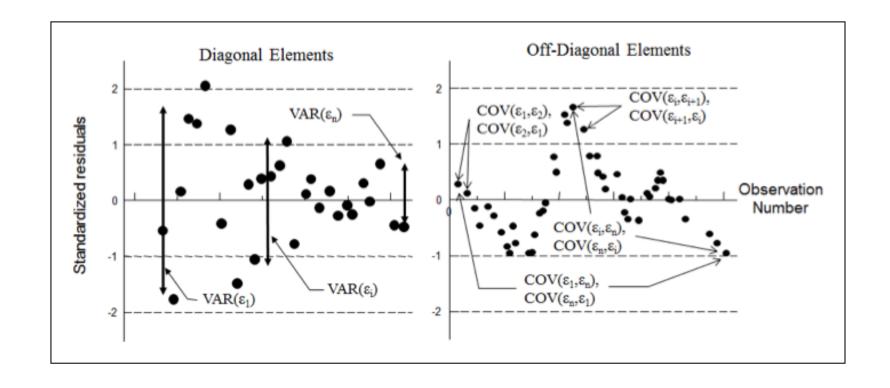
$$\hat{\sigma}^2\{\mathbf{e}\} = \begin{bmatrix} 10.404782 & -3.213241 & -2.907218 & -3.213241 & -1.071080 \\ -3.213241 & 9.410207 & -3.213241 & -3.901793 & 0.918069 \\ -2.907218 & -3.213241 & 10.404782 & -3.213241 & -1.071080 \\ -3.213241 & -3.901793 & -3.213241 & 9.410207 & 0.918069 \\ -1.071080 & 0.918069 & -1.071080 & 0.918069 & 0.306023 \end{bmatrix}$$

The **variances in residuals** (shown in the figure on the left) correspond to the diagonal elements of the variance-covariance matrix



$$\hat{\sigma}^2\{\mathbf{e}\} = \begin{bmatrix} 10.404782 & -3.213241 & -2.907218 & -3.213241 & -1.071080 \\ -3.213241 & 9.410207 & -3.213241 & -3.901793 & 0.918069 \\ -2.907218 & -3.213241 & 10.404782 & -3.213241 & -1.071080 \\ -3.213241 & -3.901793 & -3.213241 & 9.410207 & 0.918069 \\ -1.071080 & 0.918069 & -1.071080 & 0.918069 & 0.306023 \end{bmatrix}$$

The **covariances in residuals** (shown in the figure on the right) correspond to the off-diagonal elements of the variance-covariance matrix



The estimated variances are oftentimes substantially different so the magnitude of each residual is considered relative to its standard error

$$r_i = \frac{e_i}{s\{e_i\}}$$
 where $s\{e_i\} = \sqrt{\text{MSE}(1 - h_{ii})}$

These are referred to as **studentized residuals**.

For example,
$$r_1 = \frac{-0.08149425}{\sqrt{10.404782}} = -0.02526449$$

```
# Compute the residual standard errors for all observations
> se = sqrt(diag(MSE * (1 - h)))
3.2256444 3.0676061 3.2256444 3.0676061 0.5531934
# Compute studentized residuals
> resid(lm.1) / se
-0.02526449 -1.49330194 1.29540187 0.34852517 -1.05801571
> rstandard(lm.1)
-0.02526434 -1.49329323 1.29539431 0.34852314 -1.05800954
```

The hat matrix was defined earlier as

$$\mathbf{H}_{n\times n} = \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'$$

The fitted values, the residuals, and the variance–covariance matrix of the residuals can be expressed via the hat matrix

$$\mathbf{\hat{Y}}_{n \times 1} = \mathbf{HY}$$
 $\mathbf{e}_{n \times 1} = (\mathbf{I} - \mathbf{H})\mathbf{Y}$
 $\boldsymbol{\sigma}^2 \{ \mathbf{e} \} = \sigma^2 \ (\mathbf{I} - \mathbf{H})$

The hat matrix is also useful in identifying outlying *X*-values. In particular the elements on the main diagonal

```
0.21839080
                 0.24137931 0.21839080
                                        0.24137931
                                                     0.08045977
                 0.29310345 0.24137931
     0.24137931
                                        0.29310345 -0.06896552
                 0.24137931 0.21839080
h =
     0.21839080
                                        0.24137931
                                                     0.08045977
     0.24137931
                 0.29310345 0.24137931
                                        0.29310345
                                                    -0.06896552
     0.08045977 -0.06896552 0.08045977
                                       -0.06896552
                                                    0.97701149
```

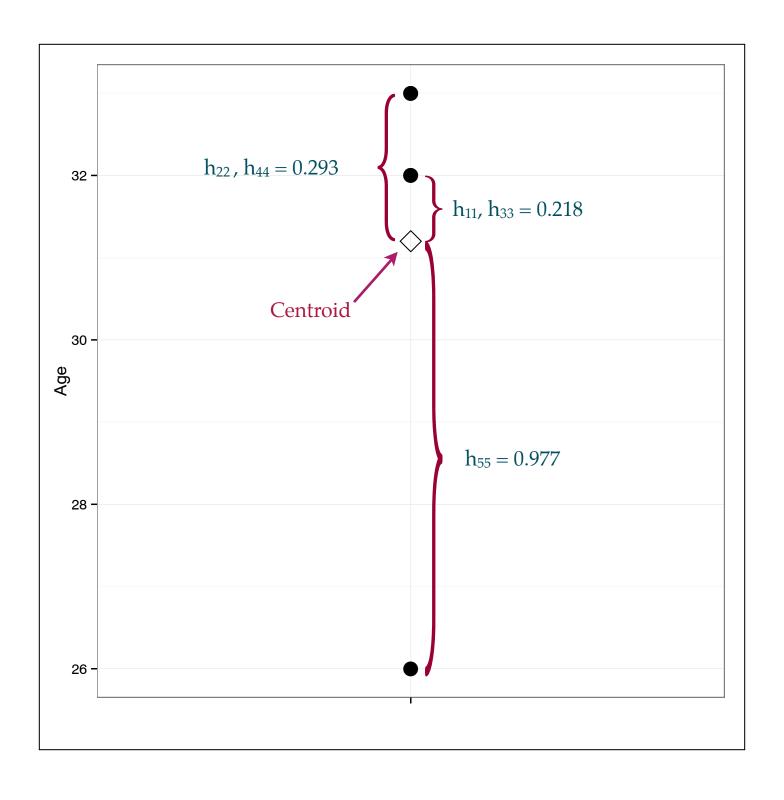
The h_{ii} elements measure the **multivariate distance** between the X-values for the ith case and the means of the X-values for all n cases (i.e., the centroid)

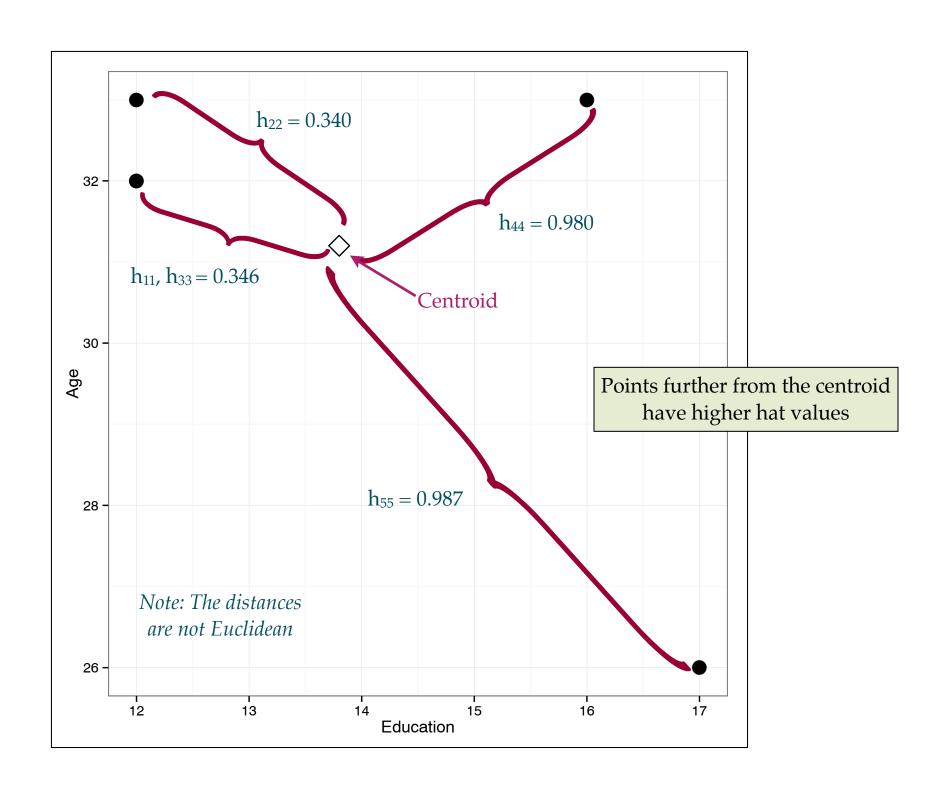
The h_{ii} elements have the following properties

$$0 \le h_{ii} \le 1$$

$$\sum_{i=1}^{n} h_{ii} = p$$

The larger the value of h_{ii} , the smaller the variance of the residual e_i , hence the closer the fitted value will tend to be to Y_i





Identifying High-Leverage Cases

The mean leverage value is

$$\bar{h} = \frac{\sum_{i=1}^{n} h_{ii}}{n} = \frac{p}{n}$$

Leverage values over *twice* as *large* as the mean leverage are considered **outlying cases** with regard to their *X*-values

In the data we have, the **mean leverage value is 0.4**. Any observation with a *hat value* ≥ 0.8 would be considered an outlying case (high leverage) based on its *X*-values (in our case observations 4 and 5)

```
# Obtain the hat-values (leverage) values
> hatvalues(lm.1)
0.2183908 0.2931034 0.2183908 0.2931034 0.9770115
# Mean hat value
> mhat = mean(hatvalues(lm.1))
> mhat
[1] 0.4
# Mean hat-value
> mean(hatvalues(lm.1))
[1] 0.4
# Is hat-value > mean hat-value
> hatvalues(lm.1) > mhat
FALSE FALSE FALSE TRUE
```

We can obtain regression results from just a few summary results, namely the correlation matrix, and standard deviations.

The standardized regression coefficients can be computed using

$$\hat{\boldsymbol{\beta}} = (\mathbf{r}_{\mathbf{X}\mathbf{X}})^{-1}\mathbf{r}_{\mathbf{X}\mathbf{Y}}$$

To obtain the unstandardized regression coefficients

$$\mathbf{b} = \hat{oldsymbol{eta}} rac{\hat{\sigma}_Y}{\hat{\sigma}_i}$$

We can obtain the multiple R² using

$$R^2 = \sum \hat{\beta} \, \mathbf{r}_{\mathbf{X}\mathbf{Y}}$$

```
# Obtain the correlation matrix
> myCor = cor(myData[ , c("wage", "age", "educ")])
> myCor
                                  educ
                       age
           wage
wage 1.0000000 0.4225006 -0.2314938
age 0.4225006 1.0000000 -0.6399446
educ -0.2314938 -0.6399446 1.0000000
# Matrix of r(XY)
> rXY = myCor[2:3, 1]
# Matrix of r(XX)
> rXX = myCor[2:3, 2:3]
# Compute regression coefficients
> b = solve(rXX) %*% rXY
> b
                         These are the regression coefficients for the model
           \lceil,1\rceil
age 0.46464173
                                    z.wage ~ z.age + z.educ
educ 0.06585118
```

```
# Obtain R^2
> sum(rXY * b)
[1] 0.1810673
# Obtain the SDs for the outcome and predictors
> s = apply(myData[c("wage", "age", "educ")], MARGIN = 2, FUN = sd)
# Separate the SDs
> sY = s[1]
> sX = s[2:3]
# Compute unstandardized regression coefficients
> (sY / sX) * b
          [,1]
                      These are the unstandardized regression coefficients from
age 0.5491758
                                  the model wage ~ age + educ
educ 0.0921978
```

Multicollinearity

Several problems when predictors are highly correlated

- Adding/deleting a predictor changes the regression coefficients
- Additional SS associated with predictor variable varies depending on which variables are in model
- Estimated SEs become large
- Tests for the estimated coefficients may not be statistically significant even though the test for the model is

Indicators of multicollinearity include

- Large changes in estimated coefficient when a predictor is added or removed
- Nonsignificant results for important predictors (based on theory or prior experience)
- Estimated coefficients with an unexpected sign (based on theory or prior experience)
- Wide CIs for the regression coefficients that represent important predictors (based on theory or prior experience)

Variance Inflation Factor (VIF)

Variance inflation factors measure how much the variances of the estimated regression coefficients are inflated as compared to when the predictors are not linearly related (i.e., independent)

The precision of the OLS estimated regression coefficients as measured by their variances is

$$\sigma^2 \{ \mathbf{b} \} = \sigma^2 \ (\mathbf{X}' \mathbf{X})^{-1}$$

For purposes of measuring the impact of multicollinearity, it is **useful to work with the standardized regression model**. In this model, the **X'X** matrix is just the correlation matrix of the *X*-variables

The variances for the standardized coefficients are found by taking the k^{th} diagonal element of the inverse of the correlation matrix

```
# Fit a model with multiple predictors
> lm.2 = lm(wage \sim age + educ + status, data = myData)
# Get the design matrix
> X = model.matrix(lm.2)
# Inverse of the correlation matrix for the predictors
# Omit the column of ones
> solve(cor(X[ , -1]))
                 age educ statusSingle
   3.281682 1.6343002 1.8150271
age
educ 1.634300 1.8843962 0.6291698
statusSingle 1.815027 0.6291698 2.0743555
> vif(lm.2)
    age educ status
3.281682 1.884396 2.074355
```

The diagonal element $(VIF)_k$ is called the variance inflation factor for b'_k and is equal to

$$(VIF)_k = (1 - R_k^2)^{-1}$$
 where $k = 1, 2, \dots, (p - 1)$

where R^2_k is the coefficient of multiple determination when x_k is regressed on the p-2 other X variables in the model.

The VIF = 1 when $R^2_k = 0$ (x_k is not related to the other predictors)

The VIF > 1 when $R^2_k \neq 0$ which indicates an inflated variance for b'_k (i.e., t-values too small; CIs too large)

The largest VIF is often used as a diagnostic indicator of the severity of multicollinearity. A maximum VIF > 10 is usually considered problematic and is unduly influencing the OLS estimates

The mean VIF also provides information about multicollinearity, where the mean VIF is

$$\overline{\text{VIF}} = \frac{\sum_{k=1}^{p-1} \text{VIF}_k}{p-1}$$

Mean VIFs > 1 also indicate model problems due to multicollinearity