Assessing Accuracy

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When we assume f is linear, we say

$$Y = f(X) + \epsilon = \beta_0 + \beta_1 X + \epsilon$$

- \triangleright where β_0 is the intercept term.
 - ightharpoonup This is the expected value of Y when X=0.
- ightharpoonup and β_1 is the slope.
 - ▶ This is the average increase in Y for a one-unit increase in X.

The model

$$Y = \beta_0 + \beta_1 X + \epsilon$$

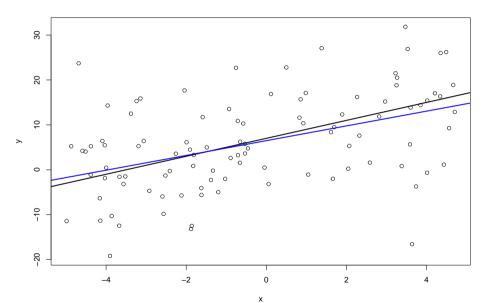
defines the (unknown) population regression line, the best linear approximation to the true relationship between X and Y.

The estimated line

$$\hat{y} = \hat{\beta}_0 + \hat{\beta}_1 x$$

is the *least squares regression line*.

```
f.x <- function(x){2*x + 7 + rnorm(length(x),0,10)}
x <- runif(100, -5, 5)
y <- f.x(x)
plot(x,y)
abline(7, 2, col='black', lwd=2)
abline(lm(y~x), col='blue', lwd=2)</pre>
```



Example: Generating Many Samples

```
rand.lines <- function(){</pre>
  x \leftarrow runif(100, -5, 5)
  y \leftarrow 2*x + 7 + rnorm(length(x), 0, 10)
  lm(y ~ x)$coefficients
coefs <- replicate(25, rand.lines())</pre>
colfunc <- colorRampPalette(c("red","yellow","springgreen","royalblue"))</pre>
colrs <- colfunc(25)
plot(-5:5, 2*(-5:5)+7, type='l', xlab='x', ylab='y')
for(i in 1:25) abline(coefs[.i], col=colrs[i])
abline(7, 2, lwd=3)
```

Example: Generating Many Samples

```
rand.lines <- function(){
  x \leftarrow runif(100, -5, 5)
  y \leftarrow 2*x + 7 + rnorm(length(x), 0, 10)
  lm(v ~ x)$coefficients
coefs <- replicate(25, rand.lines())</pre>
colfunc <- colorRampPalette(c("red","yellow","springgreen","royalblue"))</pre>
colrs <- colfunc(25)</pre>
plot(-5:5, 2*(-5:5)+7, type='l', lwd=2, xlab='x', ylab='y')
for(i in 1:25) abline(coefs[,i], col=colrs[i])
```

Least squares estimates are unbiased. Idea:

- ▶ Take a large number of samples and calculate $\hat{\beta}_0$ and $\hat{\beta}_1$ for each.
- ▶ If we were to find the mean of all the estimates of $\hat{\beta}_0$, it would be β_0 .
- ightharpoonup ... and if we were to find the mean of all the estimates of $\hat{\beta}_1$, it would be β_1 .
- ▶ We can see this visualized in the previous plot.

As in using \bar{x} to estimate μ , a regression line from a single sample may or may not be a good estimate.

- ► How variable is it?
 - ▶ When we use \bar{x} to estimate μ , the variability is

$$Var(\bar{x}) = SE(\bar{x})^2 = \frac{\sigma^2}{n}$$

ightharpoonup SE tells us roughly how far a typical estimate differs from μ .

So what about the regression line?

For \hat{eta}_0 ,

$$SE(\hat{\beta}_0)^2 = \sigma^2 \left[\frac{1}{n} + \frac{\bar{x}^2}{\sum_{i=1}^n (x_i - \bar{x})^2} \right]$$

and for $\hat{\beta}_1$,

$$SE(\hat{\beta}_1)^2 = \frac{\sigma^2}{\sum_{i=1}^n (x_i - \bar{x})^2}$$

where $\sigma^2 = \text{Var}(\epsilon)$.

lacktriangle Assumption: the errors ϵ_i are uncorrelated and have common variance.

Estimating σ

In general, σ is unknown, but can be estimated from the data:

$$\hat{\sigma} = \mathsf{RSE} = \sqrt{\frac{\mathsf{RSS}}{(n-2)}}$$

▶ This is also called the *residual standard error*.

Confidence Intervals for β_0 and β_1

A general confidence interval looks like

point estimate
$$\pm$$
 (critical value) \times (standard error)

For β_i ,

$$\hat{eta}_i \pm t_{df,\alpha/2} imes \mathsf{SE}(\hat{eta}_i)$$

▶ We use the t-distribution under the assumption that the errors are approximately Gaussian (normal).

The most common hypothesis test in this setting involves

- ▶ (Null hypothesis) H_0 : There is no relationship between X and Y.
- lacktriangle (Alternative hypothesis) H_A : There is some relationship between X and Y.

Mathematically, this is just

$$H_0: \beta_1=0$$

versus

$$H_A: \beta_1 \neq 0$$

Because, if $\beta_1=0$, then the model is just $Y=\beta_0+\epsilon$, which does not depend on X.

Note: in the model $Y = \beta_0 + \epsilon$, we find $\hat{\beta}_0 = \bar{y}$.

Two ways to test these hypotheses:

- 1. Use the confidence interval approach (check if 0 is in the interval for $\hat{\beta}_1$).
- 2. Compute a test statistic

$$t = \frac{\hat{\beta}_1 - 0}{\mathsf{SE}(\hat{\beta}_1)}$$

which measures how many standard deviations $\hat{\beta}_1$ is from 0.

From here, we typically calculate the *p-value*, or the probability of observing a value as extreme as $\hat{\beta}_1$ if in fact $\beta_1 = 0$.

In practice, we never do this by hand.

```
mod1 <- lm(Loblolly$age ~ Loblolly$height)
summary(mod1)
##
## Call:
## lm(formula = Loblolly$age ~ Loblolly$height)
##
## Residuals:
##
      Min
               10 Median
                              30
                                     Max
## -2.5528 -0.7378 0.1421 0.6925 2.8966
##
## Coefficients:
##
                  Estimate Std. Error t value Pr(>|t|)
  (Intercept) 0.757380 0.229203 3.304 0.00141 **
## Loblolly$height 0.378274 0.005979 63.272 < 2e-16 ***
```

Assessing Model Accuracy

Having concluded that β_1 is nonzero, we want to examine the extent to which the model fits the data.

Linear regression model quality assessed using two measures:

- 1. Residual standard error
- 2. R^2

Residual standard error

Recall: RSE = $\hat{\sigma}$.

- ► This is a measure of how far on average linear regression line estimates deviate from the truth.
 - ► A "good" RSE will depend on problem context (e.g., units).
- RSE is considered a lack of fit measure.
 - If predictions are very close to true outcomes, RSE will be small (and vice versa).

R^2 Statistic

RSE is measured in units of Y, so it may be unclear what a "good" RSE is.

The R^2 statistic

- is the proportion of variance explained by the model.
- ▶ always takes values between 0 and 1.

$$R^2 = \frac{\mathsf{TSS} - \mathsf{RSS}}{\mathsf{TSS}} = 1 - \frac{\mathsf{RSS}}{\mathsf{TSS}}$$

where
$$TSS = \sum (y_i - \bar{y})^2$$

Sum of Squares

- ▶ TSS is the *total sum of squares*, the total variance in *Y*.
- ▶ RSS is the *residual sum of squares*, the variability leftover after the regression is performed.
- ▶ Another measure, ESS, is the *explained sum of squares* and is the variability in *Y* that is explained by the regression model:

$$TSS = RSS + ESS$$

Thus, $R^2 = \frac{\text{ESS}}{\text{TSS}}$ is the proportion of variability in Y that can be explained by the linear regression model.

R^2 Statistic

"Good" R^2 values are those closer to 1.

... How close to 1?

It depends!

Correlation

We can also measure the (linear) correlation between two variables.

$$Cor(X,Y) = R = \frac{\sum_{i=1}^{n} (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_{i=1}^{n} (x_i - \bar{x})^2} \sqrt{\sum_{i=1}^{n} (y_i - \bar{y})^2}}$$

In the linear regression context, the square of the correlation is the R^2 we just saw.

Overall model fit

[1] 0.9899132

```
mod1 <- lm(Loblolly$age ~ Loblolly$height)
anova (mod1)
## Analysis of Variance Table
##
## Response: Loblolly$age
##
                 Df Sum Sq Mean Sq F value Pr(>F)
## Loblolly$height 1 5076 5076.0 4003.3 < 2.2e-16 ***
## Residuals 82 104 1.3
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
cor(Loblolly$height, Loblolly$age)
```