R01 - Simple linear regression

STAT 5870 (Engineering) Iowa State University

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Telomere length

http://www.pnas.org/content/101/49/17312

People who are stressed over long periods tend to look haggard, and it is commonly thought that psychological stress leads to premature aging [as measured by decreased telomere length]

examine the importance of ... caregiving stress (...number of years since a child's diagnosis [of a chronic disease]) [on telomere length]

Telomere length values were measured from DNA by a quantitative PCR assay that determines the relative ratio of telomere repeat copy number to single-copy gene copy number (T/S ratio) in experimental samples as compared with a reference DNA sample.

Data



Data with regression line



Simple Linear Regression

The simple linear regression model is

```
Y_i \stackrel{ind}{\sim} N(\beta_0 + \beta_1 X_i, \sigma^2)
```

where Y_i and X_i are the response and explanatory variable, respectively, for individual *i*.

Terminology (all of these are equivalent):

response	explanatory
outcome	covariate
dependent	independent
endogenous	exogenous

Model

Simple linear regression - visualized Simple linear regression model



Explanatory variable

Parameter interpretation

Recall:

$$E[Y_i|X_i = x] = \beta_0 + \beta_1 x \qquad Var[Y_i|X_i = x] = \sigma^2$$

• If $X_i = 0$, then $E[Y_i | X_i = 0] = \beta_0$.

 β_0 is the expected response when the explanatory variable is zero.

• If X_i increases from x to x + 1, then

 β_1 is the expected increase in the response for each unit increase in the explanatory variable.

• σ is the standard deviation of the response for a fixed value of the explanatory variable.

Simple linear regression - visualized



Errors v residuals

Remove the mean:

$$Y_i = \beta_0 + \beta_1 X_i + e_i \qquad e_i \stackrel{iid}{\sim} N(0, \sigma^2)$$

So the error is

$$e_i = Y_i - (\beta_0 + \beta_1 X_i)$$

which we approximate by the residual

$$r_i = \hat{e}_i = Y_i - (\hat{\beta}_0 + \hat{\beta}_1 X_i)$$

These residuals we will use for a number of purposes including

- assessing model assumptions,
- identifying outliers, and
- estimating error variance.

Estimators

The least squares (minimize $\sum_{i=1}^{n} r_i^2$), maximum likelihood, and Bayesian estimators (prior $1/\sigma^2$) are

$$\hat{\beta}_{1} = SXY/SXX$$

$$\hat{\beta}_{0} = \overline{Y} - \hat{\beta}_{1}\overline{X}$$

$$\hat{\sigma}^{2} = SSE/(n-2) \qquad df = n-2$$

$$\overline{X} = \frac{1}{n}\sum_{i=1}^{n}X_{i}$$

$$\overline{Y} = \frac{1}{n}\sum_{i=1}^{n}Y_{i}$$

$$SXY = \sum_{i=1}^{n}(X_{i} - \overline{X})(Y_{i} - \overline{Y})$$

$$SXX = \sum_{i=1}^{n}(X_{i} - \overline{X})(X_{i} - \overline{X}) = \sum_{i=1}^{n}(X_{i} - \overline{X})^{2}$$

$$SSE = \sum_{i=1}^{n}r_{i}^{2}$$

Residuals



Residuals



How certain are we about $\hat{\beta}_0$ and $\hat{\beta}_1$?

We quantify this uncertainty using their standard errors (or posterior scale parameters):

$$SE(\hat{\beta}_0) = \hat{\sigma}\sqrt{\frac{1}{n} + \frac{\overline{X}^2}{(n-1)s_X^2}} \qquad df = n-2$$

$$SE(\hat{\beta}_1) = \hat{\sigma}\sqrt{\frac{1}{(n-1)s_X^2}} \qquad df = n-2$$

$$\begin{array}{rl} s_X^2 &= SXX/(n-1)\\ s_Y^2 &= SYY/(n-1)\\ SYY &= \sum_{i=1}^n (Y_i - \overline{Y})^2 \end{array}$$

$$\begin{array}{ll} r_{XY} &= \frac{SXY/(n-1)}{s_X s_Y} & \mbox{corr}\\ R^2 &= r_{XY}^2 = \frac{SST-SSE}{SST} & \mbox{corr}\\ SST &= SYY = \sum_{i=1}^n (Y_i - \overline{Y})^2 \end{array}$$

correlation coefficient coefficient of determination

The coefficient of determination (R^2) is the proportion of the total response variation explained by the model.

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Default Bayesian analysis of the simple linear regression model

If we assume the default prior $p(\beta_0,\beta_1,\sigma^2)\propto 1/\sigma^2,$ then the marginal posteriors for the mean parameters are

 $\beta_j | y \sim t_{n-2}(\hat{\beta}_j, SE(\hat{\beta}_j)^2).$

We can construct a 100(1-a)% two-sided credible interval for β_j via

$$\hat{\beta}_j \pm t_{n-2,1-a/2} SE(\hat{\beta}_j)$$

where $P(T_{n-2} < t_{n-2,1-a/2}) = 1 - a/2$ for $T_{n-2} \sim t_{n-2}$.

We can compute posterior probabilities via

$$P(\beta_j > b_j | y) = P\left(T_{n-2} > \frac{b_j - \hat{\beta}_j}{SE(\hat{\beta}_j)}\right) \quad \text{or} \quad P(\beta_j < b_j | y) = P\left(T_{n-2} < \frac{b_j - \hat{\beta}_j}{SE(\hat{\beta}_j)}\right)$$

often $b_j = 0$.

p-values and confidence interval

We can construct a 100(1-a)% two-sided confidence interval for β_j via

 $\hat{\beta}_j \pm t_{n-2,1-a/2} SE(\hat{\beta}_j).$

We can compute one-sided *p*-values, $H_0: \beta_j \ge b_j$ vs $H_A: \beta_j < b_j$ has

and $H_0: \beta_j \leq b_j$ vs $H_A: \beta_j > b_j$ has

$$p\text{-value} = P\left(T_{n-2} < \frac{\hat{\beta}_j - b_j}{SE(\hat{\beta}_j)}\right) \qquad \qquad p\text{-value} = P\left(T_{n-2} > \frac{\hat{\beta}_j - b_j}{SE(\hat{\beta}_j)}\right)$$

or two-sided p-values for $H_0: \beta_j = b_j$ vs $H_A: \beta_j \neq b_j$:

$$= 2 \times \min\left\{P\left(T_{n-2} > \frac{\hat{\beta}_j - b_j}{SE(\hat{\beta}_j)}\right), P\left(T_{n-2} < \frac{\hat{\beta}_j - b_j}{SE(\hat{\beta}_j)}\right)\right\} = 2 \times P\left(T_{n-2} < -\left|\frac{\hat{\beta}_j - b_j}{SE(\hat{\beta}_j)}\right|\right)$$

software default is usually $b_j = 0$.

by hand

Calculations "by hand" in R

```
= nrow(Telomeres)
n
Xbar = mean(Telomeres$years)
Ybar = mean(Telomeres$telomere.length)
s X = sd(Telomeres$vears)
s_Y = sd(Telomeres$telomere.length)
r_XY = cor(Telomeres$telomere.length, Telomeres$years)
SXX = (n-1)*s X^2
SYY = (n-1)*s Y^{2}
SXY = (n-1)*s X*s Y*r XY
beta1 = SXY/SXX
beta0 = Ybar - beta1 * Xbar
R_2 = r XY^2
SSE = SYY * (1-R2)
sigma2 = SSE/(n-2)
sigma = sqrt(sigma2)
SE_beta0 = sigma*sqrt(1/n + Xbar<sup>2</sup>/((n-1)*s_X<sup>2</sup>))
SE beta1 = sigma*sqrt(
                                 1/((n-1)*s_{2})
```

Calculations "by hand" in R (continued)

```
# 95% CI for beta0
beta0 + c(-1, 1) * qt(.975, df = n-2) * SE_beta0
[1] 1.251761 1.483603
# 95% CI for beta1
beta1 + c(-1, 1) * qt(.975, df = n-2) * SE_beta1
[1] -0.044785794 -0.007962836
# pualue for HO: beta0 <= 0 and P(beta0 > 0 | y)
pt(beta0 / SE_beta0, df = n - 2)
[1] 1
# pualue for HO: beta1 <= 0 and P(beta1 > 0 | y)
pt(beta1 / SE beta1, df = n - 2)
[1] 0.003102353
# pvalue for HO: beta1 = 0
2 * pt(-abs(beta1 / SE_beta1), df = n - 2)
```

[1] 0.006204706

Calculations by hand

$$\begin{array}{ll} SXX&=(n-1)s_{\tilde{X}}^2=(39-1)\times 2.9354274^2=327.4358974\\ SYY&=(n-1)s_{\tilde{Y}}^2=(39-1)\times 0.1797731^2=1.2280974\\ SXY&=(n-1)s_Xs_Yr_{XY}=(39-1)\times 2.9354274\times 0.1797731\times -0.4306534=-8.6358974\\ \hat{\beta}_1&=SXY/SXX=-8.6358974/327.4358974=-0.0263743\\ \hat{\beta}_0&=\overline{Y}-\hat{\beta}_1\overline{X}=1.2202564-(-0.0263743)\times 5.5897436=1.3676821\\ R^2&=r_{XY}^2=(-0.4306534)^2=0.1854624\\ SSE&=SYY(1-R^2)=1.2280974(1-0.1854624)=1.0003316\\ \hat{\sigma}^2&=SSE/(n-2)=1.0003316/(39-2)=0.027036\\ \hat{\sigma}&=\sqrt{\hat{\sigma}^2}=\sqrt{0.027036}=0.1644262\\ SE(\hat{\beta}_0)&=\hat{\sigma}\sqrt{\frac{1}{n}+\frac{\overline{X}^2}{(n-1)s_x^2}}=0.1644262\sqrt{\frac{1}{39}+\frac{5.5897436^2}{(39-1)*2.9354274^2}}=0.0572111\\ SE(\hat{\beta}_1)&=\hat{\sigma}\sqrt{\frac{1}{(n-1)s_x^2}}=0.1644262\sqrt{\frac{1}{(39-1)*2.9354274^2}}=0.0090867\\ p_{H_A:\beta_0\neq 0}&=2P\left(T_{n-2}<-\left|\frac{\hat{\beta}_0}{SE(\hat{\beta}_0)}\right|\right)=2P(t_{37}<-2.9025065)=0.0062047\\ CI_{95\%\beta_0}&=\hat{\beta}_0\pm t_{n-2,1-a/2}SE(\hat{\beta}_1)\\ &=-0.0263743\pm2.0261925\times 0.0090867=(-0.0447858,-0.0079628)\\ \end{array}$$

p

Regression in R

m = lm(telomere.length ~ years, Telomeres)
summary(m)

Call: lm(formula = telomere.length ~ years, data = Telomeres)

Residuals:

Min 1Q Median 3Q Max -0.42218 -0.08537 0.02056 0.10738 0.28869

Coefficients:

Estimate Std. Error t value Pr(>|t|) (Intercept) 1.367682 0.057211 23.906 <2e-16 *** years -0.026374 0.009087 -2.903 0.0062 ** ---Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.1644 on 37 degrees of freedom Multiple R-squared: 0.1855,Adjusted R-squared: 0.1634 F-statistic: 8.425 on 1 and 37 DF, p-value: 0.006205

confint(m)

2.5 % 97.5 % (Intercept) 1.25176134 1.483602799 years -0.04478579 -0.007962836

Conclusion

Telomere ratio at the time of diagnosis of a child's chronic illness is estimated to be 1.37 with a 95% credible interval of (1.25, 1.48). For each year since diagnosis, the telomere ratio decreases on average by 0.026 with a 95% credible interval of (0.008, 0.045). The proportion of variability in telomere length described by a linear regression on years since diagnosis is 18.5%.

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The correlation between chronicity of caregiving and mean telomere length is -0.445 (P < 0.01). [$R^2 = 0.198$ was shown in the plot.]

Remark I'm guessing our analysis and that reported in the paper don't match exactly due to a discrepancy in the data.

Summary

• The simple linear regression model is

$$Y_i \stackrel{ind}{\sim} N(\beta_0 + \beta_1 X_i, \sigma^2)$$

where Y_i and X_i are the response and explanatory variable, respectively, for individual i.

- Know how to use R to obtain $\hat{\beta}_0$, $\hat{\beta}_1$, $\hat{\sigma}^2$, R^2 , *p*-values, Cls, etc.
- Interpret regression output:
 - β_0 is the expected value for the response when the explanatory variable is 0.
 - β_1 is the expected increase in the response for each unit increase in the explanatory variable.
 - σ is the standard deviation of responses around their mean.
 - R^2 is the proportion of the total variation of the response variable explained by the model.