

STAT 401A - Statistical Methods for Research Workers

Simple linear regression

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last updated: October 13, 2014

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Recall the one-way ANOVA model:

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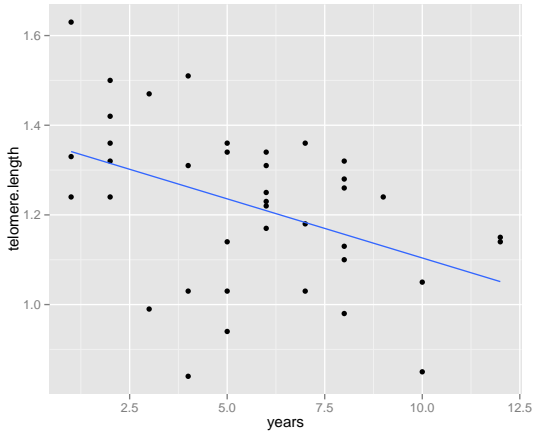
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Terminology (all of these are equivalent):

response	explanatory
outcome	covariate
dependent	independent
endogenous	exogenous



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 and the earlier onset of diseases of aging.

...

This design allowed us to examine the importance of perceived stress and measures of objective stress (caregiving status and chronicity of caregiving stress based on the number of years since a child's diagnosis).

...

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.

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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.

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$$\begin{aligned} SXY &= \sum_{i=1}^n (X_i - \bar{X})(Y_i - \bar{Y}) \\ SXX &= \sum_{i=1}^n (X_i - \bar{X})(X_i - \bar{X}) = \sum_{i=1}^n (X_i - \bar{X})^2 \\ SSE &= \sum_{i=1}^n r_i^2 \end{aligned}$$

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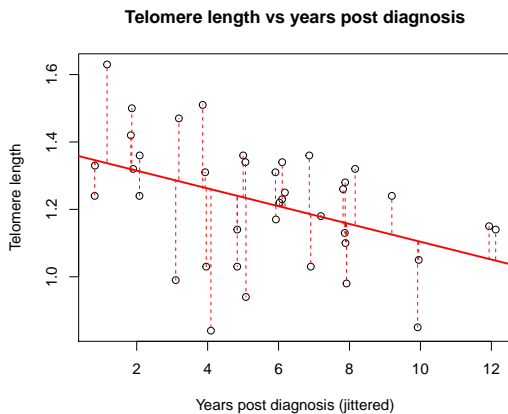
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The coefficient of determination (R^2) is the proportion of the total response variation explained by the explanatory variable(s).



Pvalues and confidence interval

We can compute two-sided pvalues via

$$2P\left(t_{n-2} < -\left|\frac{\hat{\beta}_0}{SE(\beta_0)}\right|\right) \quad \text{and} \quad 2P\left(t_{n-2} < -\left|\frac{\hat{\beta}_1}{SE(\beta_1)}\right|\right)$$

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We can construct $100(1 - \alpha)\%$ two-sided confidence intervals via

$$\hat{\beta}_0 \pm t_{n-2}(1 - \alpha/2)SE(\beta_0) \quad \text{and} \quad \hat{\beta}_1 \pm t_{n-2}(1 - \alpha/2)SE(\beta_1)$$

These provide ranges of the parameters consistent with the data.

```
DATA t;
  INFILE 'telomeres.csv' DSD FIRSTOBS=2;
  INPUT years length;
```

```
PROC CORR DATA=t;
  VAR length;
  WITH years;
  RUN;
```

The CORR Procedure

```
1 With Variables:  years
1   Variables:    length
```

Simple Statistics

Variable	N	Mean	Std Dev	Sum	Minimum	Maximum
years	39	5.58974	2.93543	218.00000	1.00000	12.00000
length	39	1.22026	0.17977	47.59000	0.84000	1.63000

Pearson Correlation Coefficients, N = 39
 Prob > |r| under H0: Rho=0

```
length
years    -0.43065
          0.0062
```

```
PROC GLM DATA=t;
  MODEL length = years / SOLUTION CLPARM;
  RUN;
```

The GLM Procedure

Number of Observations Read	39
Number of Observations Used	39

Dependent Variable: length

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	1	0.22776588	0.22776588	8.42	0.0062
Error	37	1.00033156	0.02703599		
Corrected Total	38	1.22809744			

R-Square	Coeff Var	Root MSE	length Mean
0.185462	13.47473	0.164426	1.220256

Source	DF	Type I SS	Mean Square	F Value	Pr > F
years	1	0.22776588	0.22776588	8.42	0.0062

Source	DF	Type III SS	Mean Square	F Value	Pr > F
years	1	0.22776588	0.22776588	8.42	0.0062

Parameter	Estimate	Standard Error	t Value	Pr > t	95% Confidence Limits	
Intercept	1.367682067	0.05721112	23.91	<.0001	1.251761335	1.483602799
years	-0.026374315	0.00908674	-2.90	0.0062	-0.044785794	-0.007962836

Regression in R

```
m = lm(telomere.length~years, Telomeres)
with(Telomeres, cor(telomere.length,years))
```

```
[1] -0.4307
```

```
anova(m)
```

Analysis of Variance Table

Response: telomere.length

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
years	1	0.228	0.228	8.42	0.0062 **
Residuals	37	1.000	0.027		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

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
	-0.4222	-0.0854	0.0206	0.1074	0.2887

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.36768	0.05721	23.9	<2e-16 ***
years	-0.02637	0.00909	-2.9	0.0062 **

```
---
```

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
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```
Residual standard error: 0.164 on 37 degrees of freedom
```

```
Multiple R-squared:  0.185, Adjusted R-squared:  0.163
```

```
F-statistic: 8.42 on 1 and 37 DF,  p-value: 0.0062
```

```
confint(m)
```

	2.5 %	97.5 %
(Intercept)	1.25176	1.483603
years	-0.04479	-0.007963

Conclusion

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

The zero-order correlation between chronicity of caregiving [years] and mean telomere length, r , 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.

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- Interpret SAS output
 - At a value of zero for the explanatory variable ($X_i = 0$), β_0 is the expected value for the response (Y_i).

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 - At a value of zero for the explanatory variable ($X_i = 0$), β_0 is the expected value for the response (Y_i).
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 - The coefficient of determination (R^2) is the percentage of the total response variation explained by the explanatory variable(s).