

Results

Regression output:

	Estimate	Std. Error	t value	$\Pr(> t)$
(Intercept)	9.2076	9.2999	0.99	0.3316
bioIQ	0.9014	0.0963	9.36	0.0000

Linear model:

$$\widehat{fosterIQ} = 9.2076 + 0.9014 \ bioIQ$$

$$R^2$$
:

$$R^2$$
: $R^2 = 0.78$



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Testing for the Slope - Hypotheses

➤ Is the explanatory variable a significant predictor of the response variable?

 H_0 (nothing going on):

The explanatory variable is not a significant predictor of the response variable, i.e. no relationship → slope of the relationship is 0.

 $H_0: \beta_1 = 0$

 H_A (something going on):

The explanatory variable is a significant predictor of the response variable, i.e. relationship → slope of the relationship

 $H_A: \beta_1 \neq 0$

is different than 0.



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Testing for the Slope - Mechanics

➤ Use a *t*-statistic in inference for regression

$$T = \frac{\text{point estimate - null value}}{SE}$$

$$SE_{b_1}$$

t-statistic for the slope:
$$T = \frac{b_1 - 0}{SE_{b_1}} \qquad d\!f = n-2$$



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Focus on degrees of freedom

> Degrees of freedom for linear regression:

$$>df=n-2$$

- > Lose 1 df for each parameter estimated
- ➤ In linear regression we estimate 2 parameters:

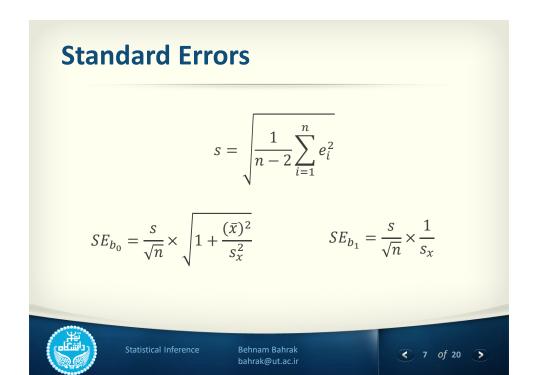
 β_0 and β_1

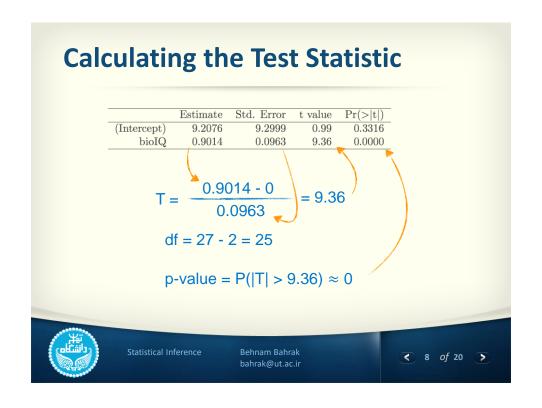


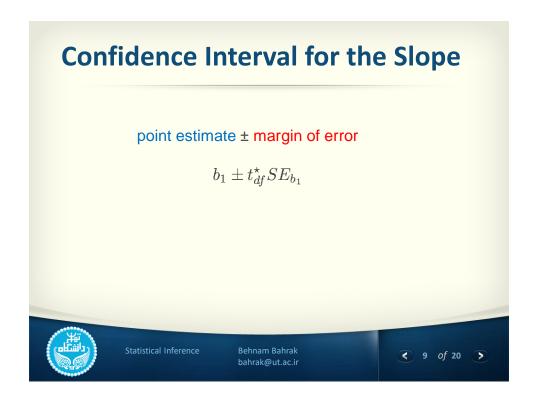
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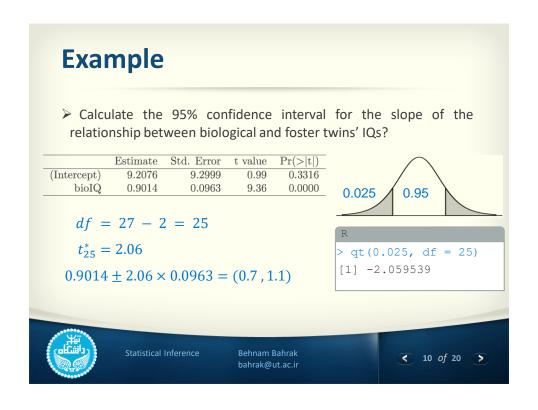
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Interpret the 95% confidence interval for the slope of the relationship between biological and foster twins' IQs:

(0.7, 1.1)

We are 95% confident that for each additional point on the biological twins' IQs, the foster twins' IQs are expected on average to be higher by 0.7 to 1.1 points.



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Recap - Inference for Regression

hypothesis test:

confidence interval:

$$T = \frac{b_1 - \text{null value}}{SE_{b_1}} \quad df = n - 2$$

$$b_1 \pm t_{df}^{\star} S E_{b_1}$$

- Null value is often 0, since we usually check for any relationship between the explanatory and the response variables.
- \triangleright Regression output gives b_1 , SE_{b_1} , and two-tailed p-value for the t-test for the slope where the null value is 0.
- Inference on the intercept is rarely done.



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Caution!

- Always be aware of the type of data you're working with: random sample, non-random sample, or population.
- > Statistical inference, and the resulting p-values, are meaningless when you already have population data.
- ➤ If you have a sample that is non-random (biased), the results will be unreliable.
- ➤ The ultimate goal is to have independent observations and you know how to check for those by now.



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Variability Partitioning

- ➤ So far: t-test as a way to evaluate the strength of evidence for a hypothesis test for the slope of relationship between x and y.
- \triangleright Alternative: consider the variability in y explained by x, compared to the unexplained variability.
- ➤ Partitioning the variability in y to explained and unexplained variability requires analysis of variance (ANOVA).



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	Df	Sum Sq	Mean Sq	F value	Pr(>F)
bioIQ	1	5231.13	5231.13	87.56	0.0000
Residuals	25	1493.53	59.74		
Total	26	6724.66			

Sum of Squares:

total variability in y: $SS_{Tot} = \sum (y - \bar{y})^2 = 6724.66$

unexplained variability in y (residuals): $SS_{Res} = \sum (y - \hat{y})^2 = \sum e_i^2 = 1493.53$

explained variability in *y*: $SS_{Reg} = 6724.66 - 1493.53 = 5231.13$



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ANOVA Output

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
bioIQ	1	5231.13	5231.13	87.56	0.0000
Residuals	25	1493.53	59.74		
Total	26	6724.66			

degrees of freedom:

total degrees of freedom: $df_{Tot} = 27 - 1 = 26$

regression degrees of freedom: $df_{Reg} = 1$ only 1 predictor

residual degrees of freedom: $df_{Res} = 26 - 1 = 25$



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	Df	Sum Sq	Mean Sq	F value	Pr(>F)
bioIQ	1	5231.13	5231.13	87.56	0.0000
Residuals	25	1493.53	59.74		
Total	26	6724.66			

mean squares

MS regression:

$$MS_{Reg} = \frac{SS_{Reg}}{df_{Reg}} = \frac{5231.13}{1} = 5231.13$$

MS residual:

$$MS_{Res} = \frac{SS_{Res}}{df_{Res}} = \frac{1493.53}{25} = 59.74$$

F statistic

ratio of explained to unexplained variability

$$F_{(1,25)} = \frac{MS_{Reg}}{MS_{Res}} = 87.56$$



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ANOVA

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
bioIQ	1	5231.13	5231.13	87.56	0.0000
Residuals	25	1493.53	59.74		
Total	26	6724.66			

 $H_0: \beta_1 = 0$

small p-value → reject H₀

 $H_A: \beta_1 \neq 0$

> The data provide convincing evidence that the slope is significantly different than 0, i.e. the explanatory variable is a significant predictor of the response variable.



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Revisiting R^2

- $ightharpoonup R^2$ is the proportion of variability in y explained by the model:
 - \triangleright large \rightarrow linear relationship between x and y exists
 - ➤ small → evidence provided by the data may not be convincing
- \triangleright Two ways to calculate R^2 :
 - (1) using correlation: square of the correlation coefficient
 - (2) from the definition: proportion of explained to total variability



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