

# Biostat 200

## Lecture 9

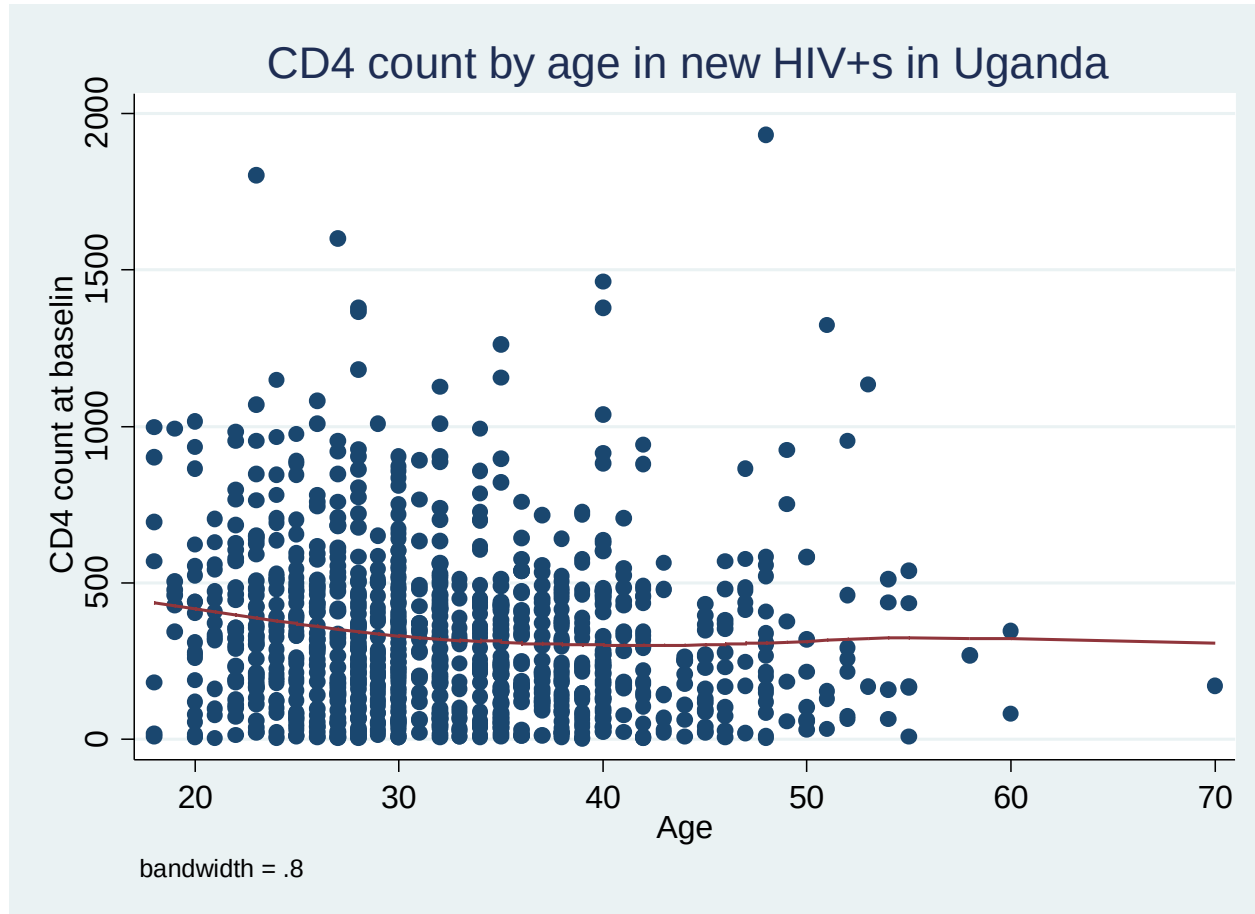
# Categorical data analyses

- Another type of paired categorical data
  - Inter-rater agreement, kappa statistic
  - Goes from 0 to 1
  - It uses expected cell frequencies, like we calculated in the previous lecture (Slides 12-13).
  - It compares observed agreement to expected agreement
    - Agreement is defined as  $\# \text{ concordant} / n$
  - In Stata, use kap command  
(<http://www.stata.com/manuals13/rkappa.pdf>)

# Scatterplot

- Back to continuous outcomes
- T-test, ANOVA, Wilcoxon rank-sum test, Kruskal-Wallis test compare 2 or more independent samples
  - e.g. CD4 cell count by sex or alcohol consumption category
- The scatterplot is a simple method to examine the relationship between 2 continuous variables

# Scatter plot



```
lowess cd4count age, xtitle(Age) ytitle(CD4 count at  
baselin) title(CD4 count by age in new HIV+s in Uganda)
```

# Correlation

- Correlation is a method to examine the relationship between 2 continuous variables
  - Does one increase with the other?
    - E.g. Does CD4 count increase with increasing age?
- Both variables are measured on the same people (or unit of analysis)
- Correlation assumes a *linear* relationship between the two variables
- Correlation is symmetric
  - The correlation of A with B is the same as the correlation of B with A

# Correlation

- Correlation is a measure of the relationship between two random variables  $X$  and  $Y$
- If the variables increase together (or oppositely), then the average of  $X*Y$  will be large (in absolute terms)
- We subtract off the mean and divide by the standard deviation to standardize (i.e. so correlations will be between -1 and 1)

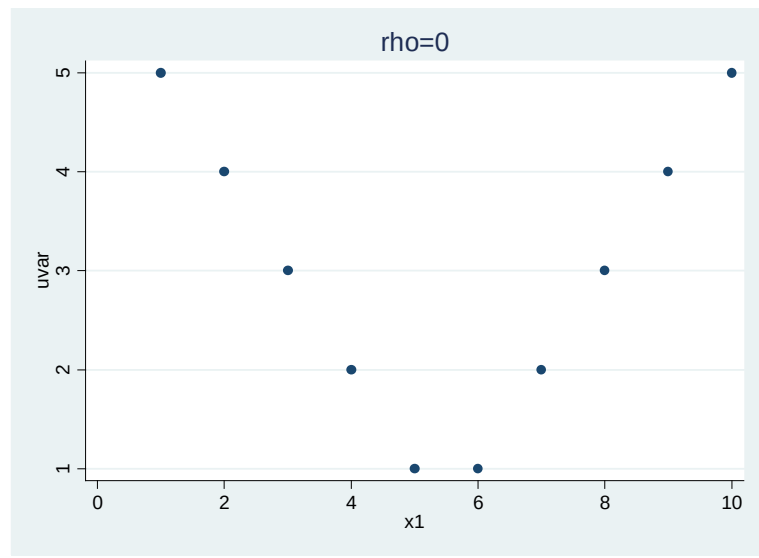
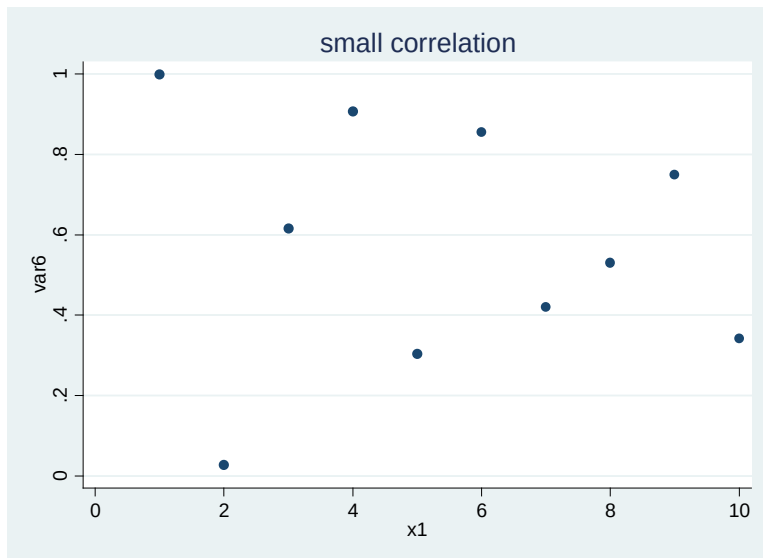
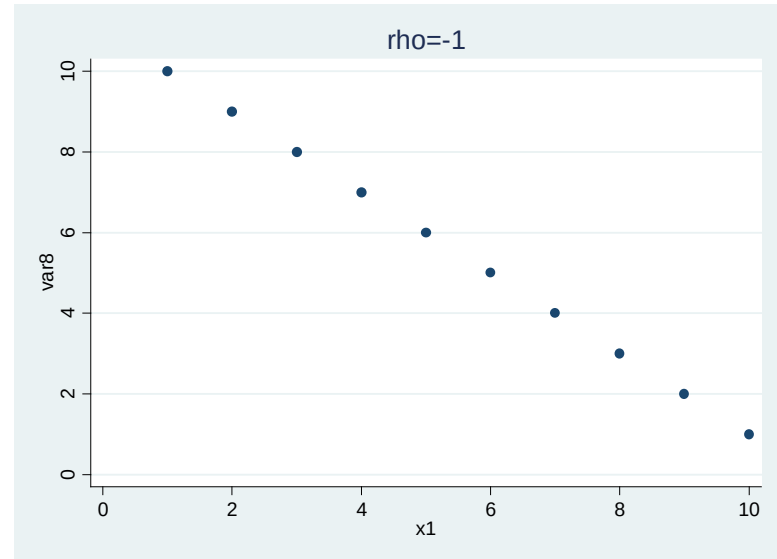
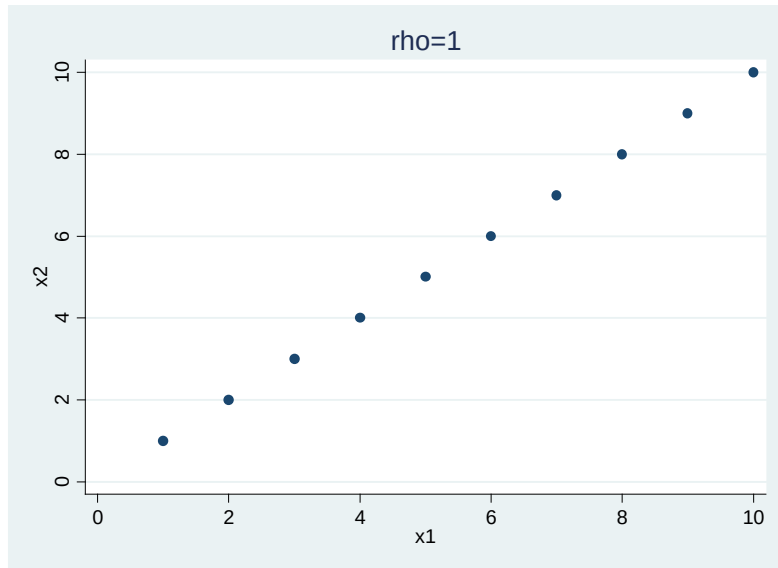
# Correlation

- A correlation is defined as

$$\rho = \text{average} \left[ \left( \frac{X - \mu_x}{\sigma_x} \right) \left( \frac{Y - \mu_y}{\sigma_y} \right) \right]$$

- Correlations can be comparable across variables with different means and variability
- Correlation does not imply causation

# Correlation





# Correlation

- $\rho$  lies between -1 and 1
- -1 and 1 are perfect correlations, 0 is no correlation
- An estimator of the population correlation  $\rho$  is Pearson's correlation coefficient denoted  $r$
- It is estimated by: 
$$r = \frac{1}{n-1} \sum_{i=1}^n \left( \frac{x_i - \bar{x}}{s_x} \right) \left( \frac{y_i - \bar{y}}{s_y} \right)$$



# Correlation

$$r = \frac{1}{n-1} \sum_{i=1}^n \left( \frac{x_i - \bar{x}}{s_x} \right) \left( \frac{y_i - \bar{y}}{s_y} \right)$$

$$= \frac{1}{(n-1)} \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\left[ \frac{\sum_{i=1}^n (x_i - \bar{x})^2}{n-1} \right]} \sqrt{\left[ \frac{\sum_{i=1}^n (y_i - \bar{y})^2}{n-1} \right]}}$$

$$= \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\left[ \sum_{i=1}^n (x_i - \bar{x})^2 \right]} \sqrt{\left[ \sum_{i=1}^n (y_i - \bar{y})^2 \right]}}$$

# Correlation: hypothesis testing

- To test whether there is a correlation between two variables, our hypotheses are

$$H_0 : \rho=0 \quad \text{and} \quad H_A : \rho \neq 0$$

- We need to calculate a test statistic for  $r$
- The test statistic is  $t = \frac{r - 0}{se(r)}$

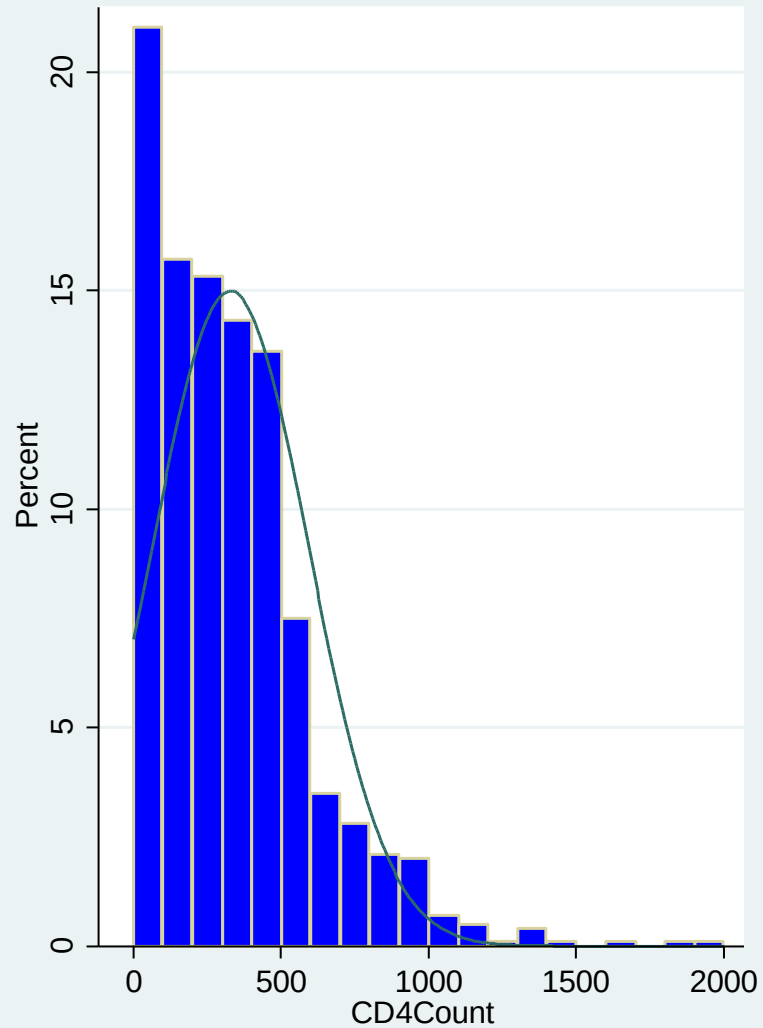
$$\text{where } se(r) = \sqrt{\frac{1 - r^2}{n - 2}}$$

$$\text{so } t = r \sqrt{\frac{n - 2}{1 - r^2}}$$

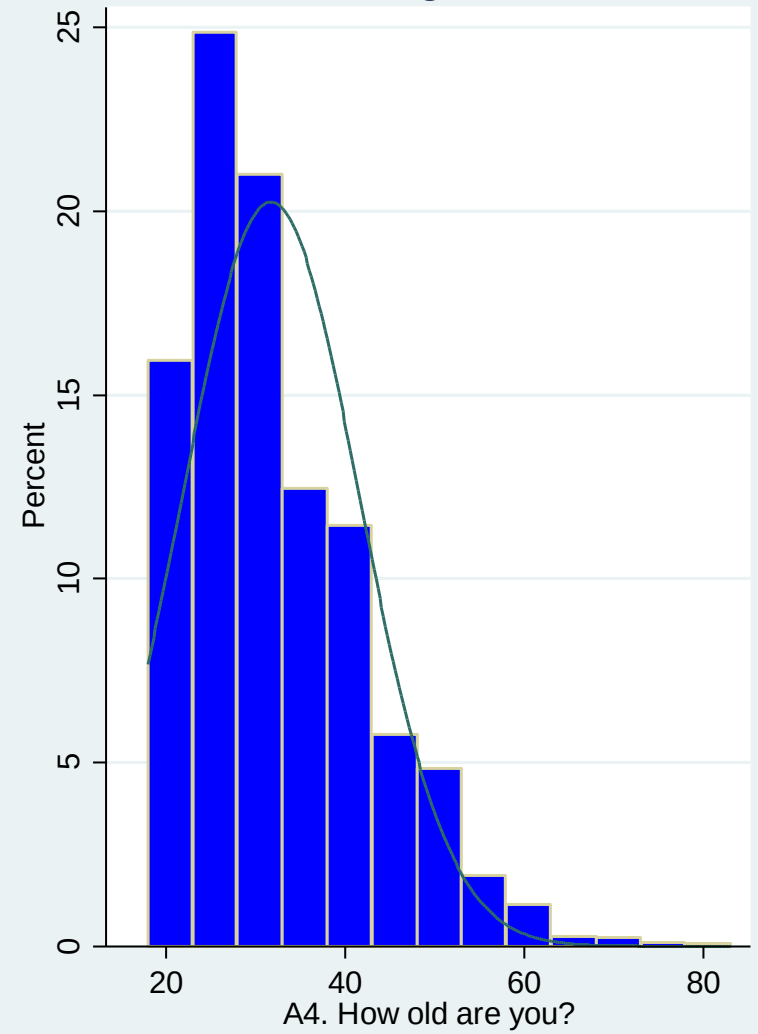
# Correlation: hypothesis testing

- The test statistic follows a t distribution with  $n-2$  degrees of freedom under the null
- The assumptions
  - The pairs of observations  $(x_i, y_i)$  were obtained from a random sample
  - $X$  and  $Y$  are normally distributed

CD4 count



Age



# Correlation example

```
corr cd4count age  
(obs=999)
```

```
              | cd4count      age_b  
-----+-----  
cd4count |      1.0000  
age_b   |     -0.1133      1.0000
```

Note that the hypothesis test is only of  $\rho=0$ , no other null  
Also note that the correlation is the linear relationship only



# Correlation example

`pwcorr var1 var2, sig obs`

```
. pwcorr cd4count age, sig obs
```

|          | cd4count                 | age_b          |
|----------|--------------------------|----------------|
| cd4count | 1.0000<br>999            |                |
| age_b    | -0.1133<br>0.0003<br>999 | 1.0000<br>3387 |

# Spearman rank correlation (nonparametric)

- Pearson's correlation coefficient is very sensitive to extreme values
- Spearman rank correlation calculates the Pearson correlation on the *ranks* of each variable
- The Pearson correlation coefficient is calculated, but the data values are replaced by the ranks
- The Spearman rank correlation coefficient is

$$r_s = \frac{1}{(n-1)} \sum_{i=1}^n \left( \frac{x_{ri} - \bar{x}_r}{s_{rx}} \right) \left( \frac{y_{ri} - \bar{y}_r}{s_{ry}} \right)$$



# Spearman rank correlation (nonparametric)

- The Spearman rank correlation ranges between -1 and 1 as does the Pearson correlation
- We can test the null hypothesis that  $\rho=0$
- The test statistic for  $n>10$  is with  $n-2$  degrees of freedom is compared to the t distribution

$$t_s = r_s \sqrt{\frac{n-2}{1-r_s^2}}$$

```
spearman  cd4count age, stats(rho obs p)
```

```
Number of obs =      999  
Spearman's rho =     -0.1319
```

```
Test of Ho: cd4count and age_b are independent  
Prob > |t| =      0.0000
```

# Matrix of Pearson correlations

- We can calculate a correlation matrix by listing multiple variable names
- Beware of which n's are used (use listwise option to get all n's equal)

```
pwcorr cd4count age nsupport_b auditc, sig obs
```

|            | cd4count                 | age_b                    | nsupport_b               | auditc         |
|------------|--------------------------|--------------------------|--------------------------|----------------|
| cd4count   | 1.0000<br>999            |                          |                          |                |
| age_b      | -0.1133<br>0.0003<br>999 | 1.0000<br>3387           |                          |                |
| nsupport_b | 0.0067<br>0.8340<br>994  | 0.3597<br>0.0000<br>3370 | 1.0000<br>3372           |                |
| auditc     | 0.0670<br>0.0392<br>948  | 0.0580<br>0.0009<br>3242 | 0.0468<br>0.0078<br>3230 | 1.0000<br>3244 |

# Matrix of Spearman correlations

spearman cd4count age nsupport\_b auditc, pw stats(rho obs p)

|               |
|---------------|
| Key           |
| rho           |
| Number of obs |
| Sig. level    |

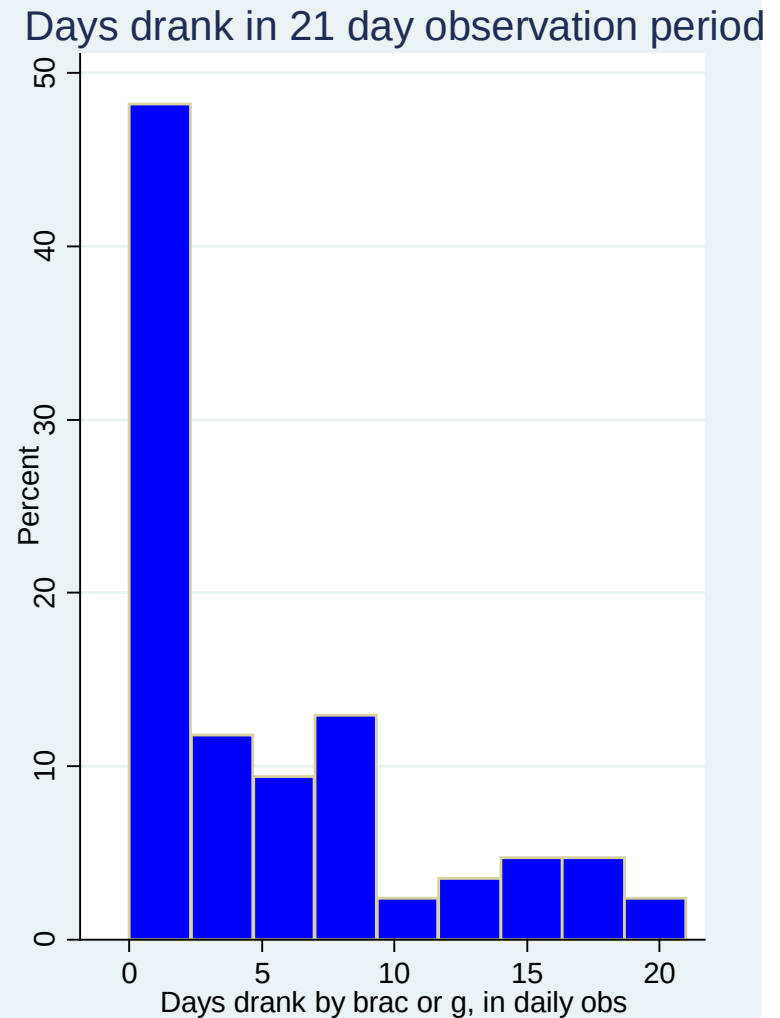
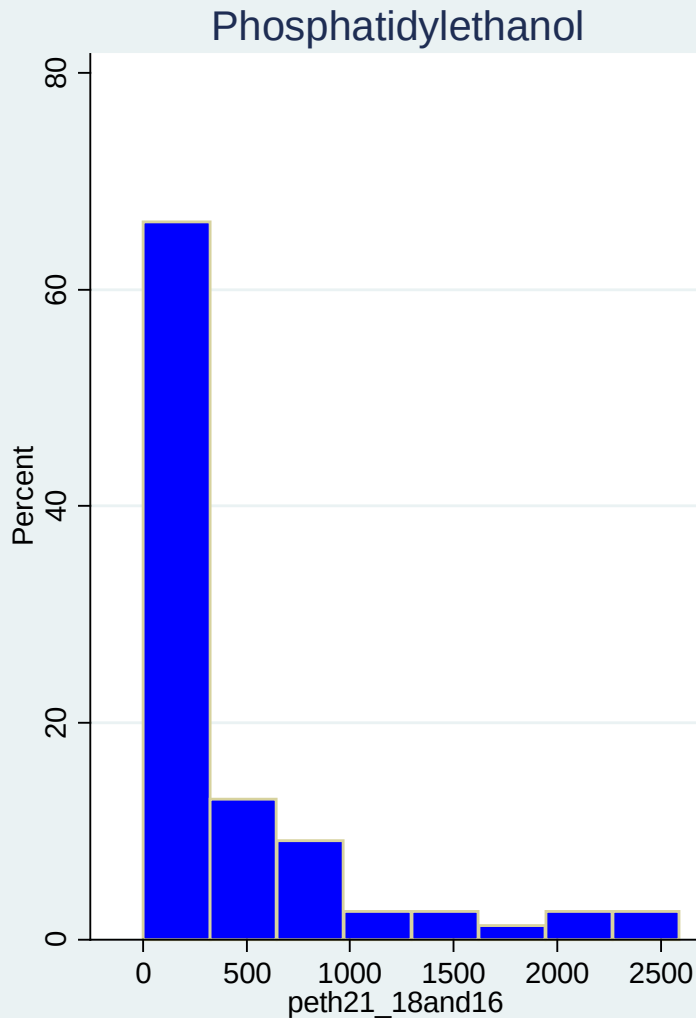
Here if you drop the “pw” option you get all n’s equal

|            | cd4count                 | age_b                    | nsuppo~b                 | auditc         |
|------------|--------------------------|--------------------------|--------------------------|----------------|
| cd4count   | 1.0000<br>999            |                          |                          |                |
| age_b      | -0.1319<br>999<br>0.0000 | 1.0000<br>3387           |                          |                |
| nsupport_b | 0.0118<br>994<br>0.7093  | 0.4055<br>3370<br>0.0000 | 1.0000<br>3372           |                |
| auditc     | 0.0945<br>948<br>0.0036  | 0.0777<br>3242<br>0.0000 | 0.0447<br>3230<br>0.0111 | 1.0000<br>3244 |

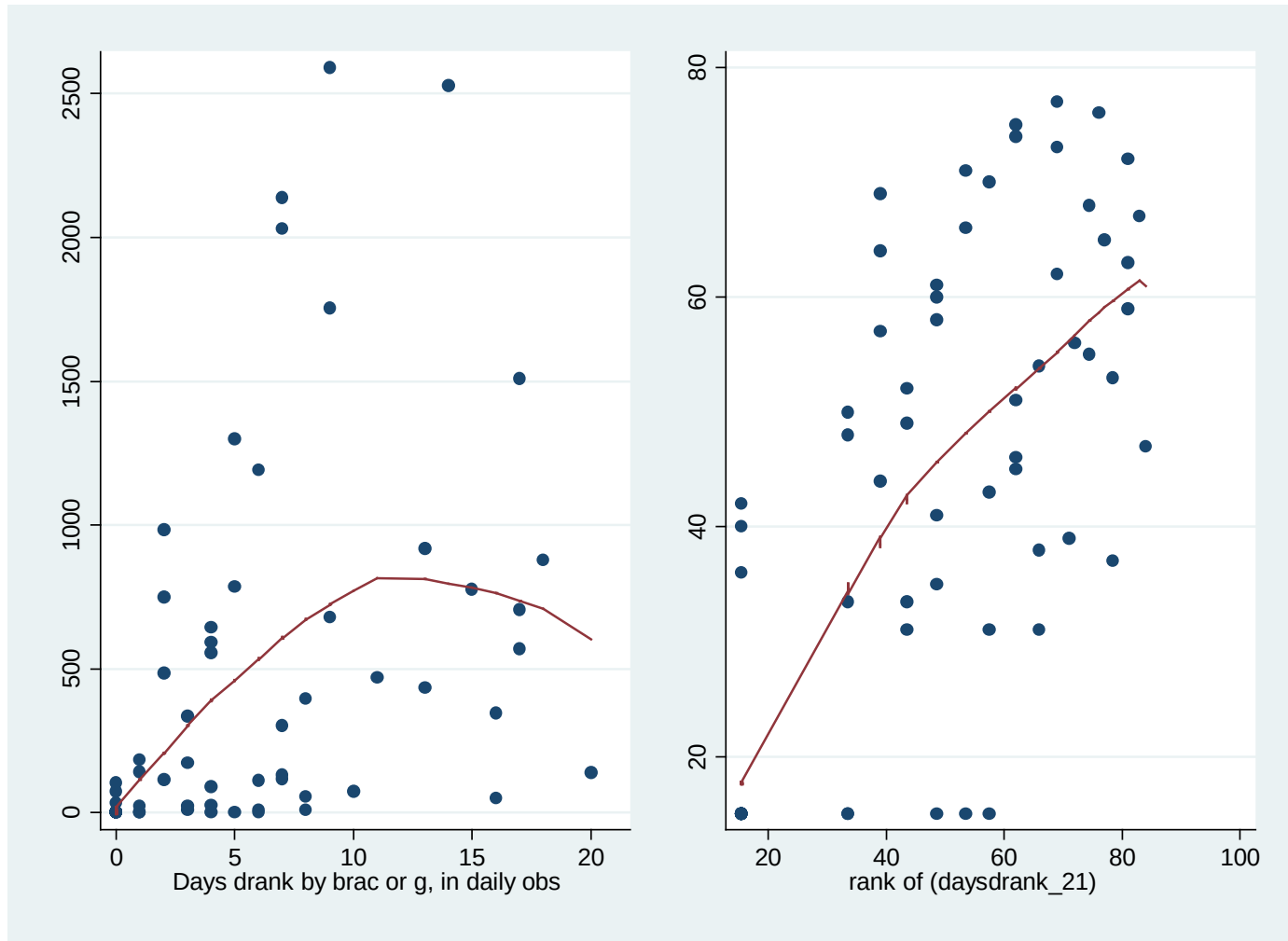
# Pearson vs. Spearman

|                                             | Pearson        | Spearman       |
|---------------------------------------------|----------------|----------------|
| Sensitivity to outliers                     | more sensitive | less sensitive |
| Can be used with ordinal data               | no             | yes            |
| Assumes data are from a normal distribution | yes            | no             |
| Detects non-linear relationship             | no             | yes            |

# Biomarker and alcohol consumption



# PEth Biomarker of alcohol consumption vs. days drinking (raw data vs. ranks)



# Pearson and Spearman correlations

```
. pwcorr peth21_18and16 daysdrank_21, obs sig
```

|              | peth2~16               | days~_21     |
|--------------|------------------------|--------------|
| peth21_18~16 | 1.0000                 |              |
| daysdrank_21 | 0.4717<br>0.0000<br>77 | 1.0000<br>85 |

```
. spearman peth21_18and16 daysdrank_21
```

Number of obs = 77  
Spearman's rho = 0.7413

Test of Ho: peth21\_18and16 and daysdrank\_21 are independent  
Prob > |t| = 0.0000

# Simple linear regression

- Correlation allows us to quantify a linear relationship between two variables
- Regression allows us to additionally estimate how a change in a random variable  $X$  corresponds to a change in random variable  $Y$

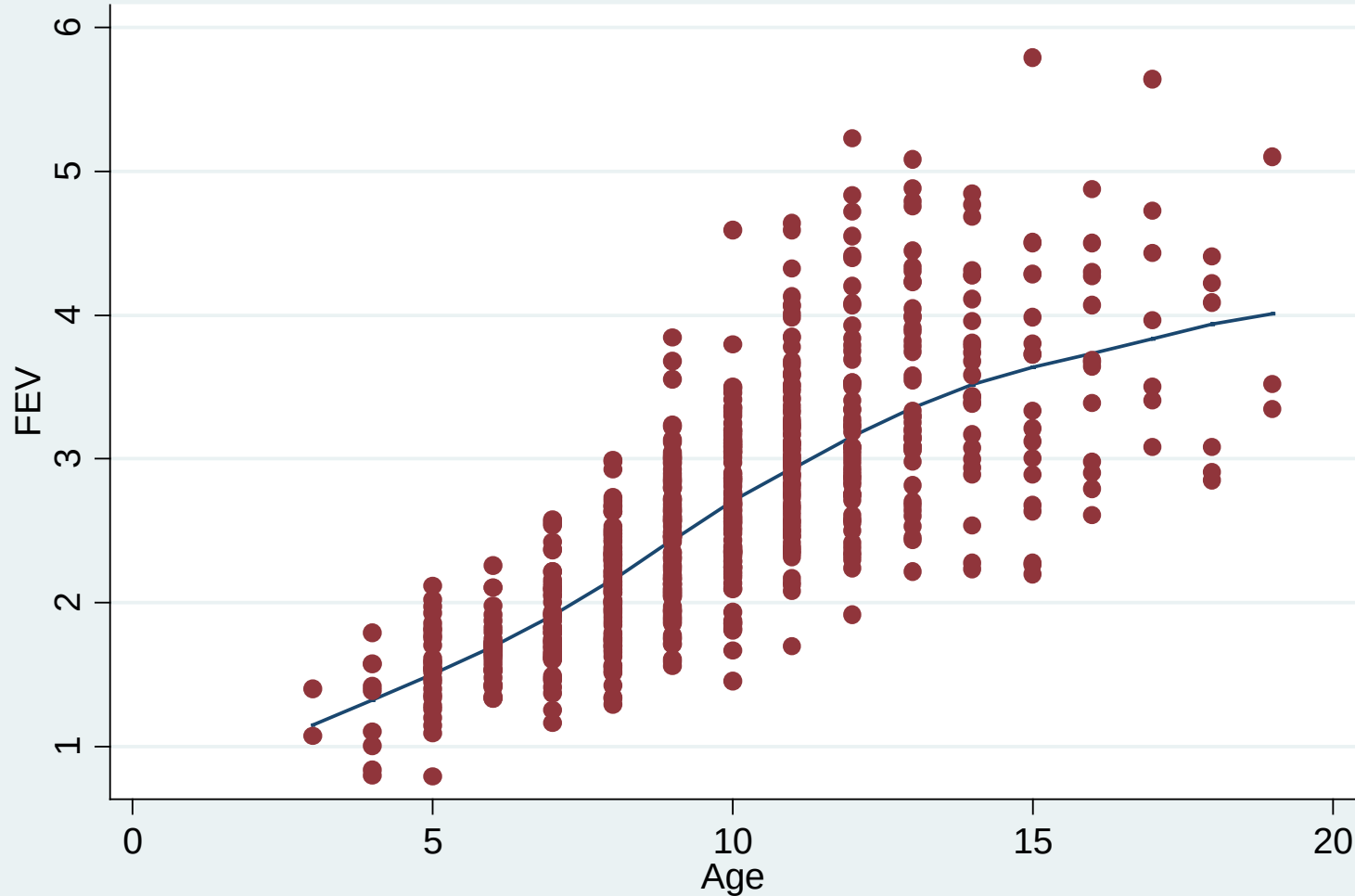


# Forced expiratory volume (FEV)

- Studies in the 1970's of children and adolescent's pulmonary function, examining their own smoking and secondhand smoke
  - FEV is the amount of air expelled in the first second of exhalation; forced expiratory volume
  - The data are cross-sectional data from a larger prospective study
- 
- Tager, I., Weiss, S., Munoz, A., Rosner, B., and Speizer, F. (1983), "Longitudinal Study of the Effects of Maternal Smoking on Pulmonary Function," *New England Journal of Medicine*, 309(12), 699-703.
  - Tager, I., Weiss, S., Rosner, B., and Speizer, F. (1979), "Effect of Parental Cigarette Smoking on the Pulmonary Function of Children," *American Journal of Epidemiology*, 110(1), 15-26.



## FEV vs age in children and adolescents



```
twoway (lowess fev age, bwidth(0.8)) (scatter fev age,
sort), ytitle(FEV) xtitle(Age) legend(off) title(FEV vs
age in children and adolescents)
```

# Correlation

```
. pwcorr fev age, sig obs
```

|     | fev    | age    |
|-----|--------|--------|
| fev | 1.0000 |        |
|     | 654    |        |
| age | 0.7565 | 1.0000 |
|     | 0.0000 |        |
|     | 654    | 654    |

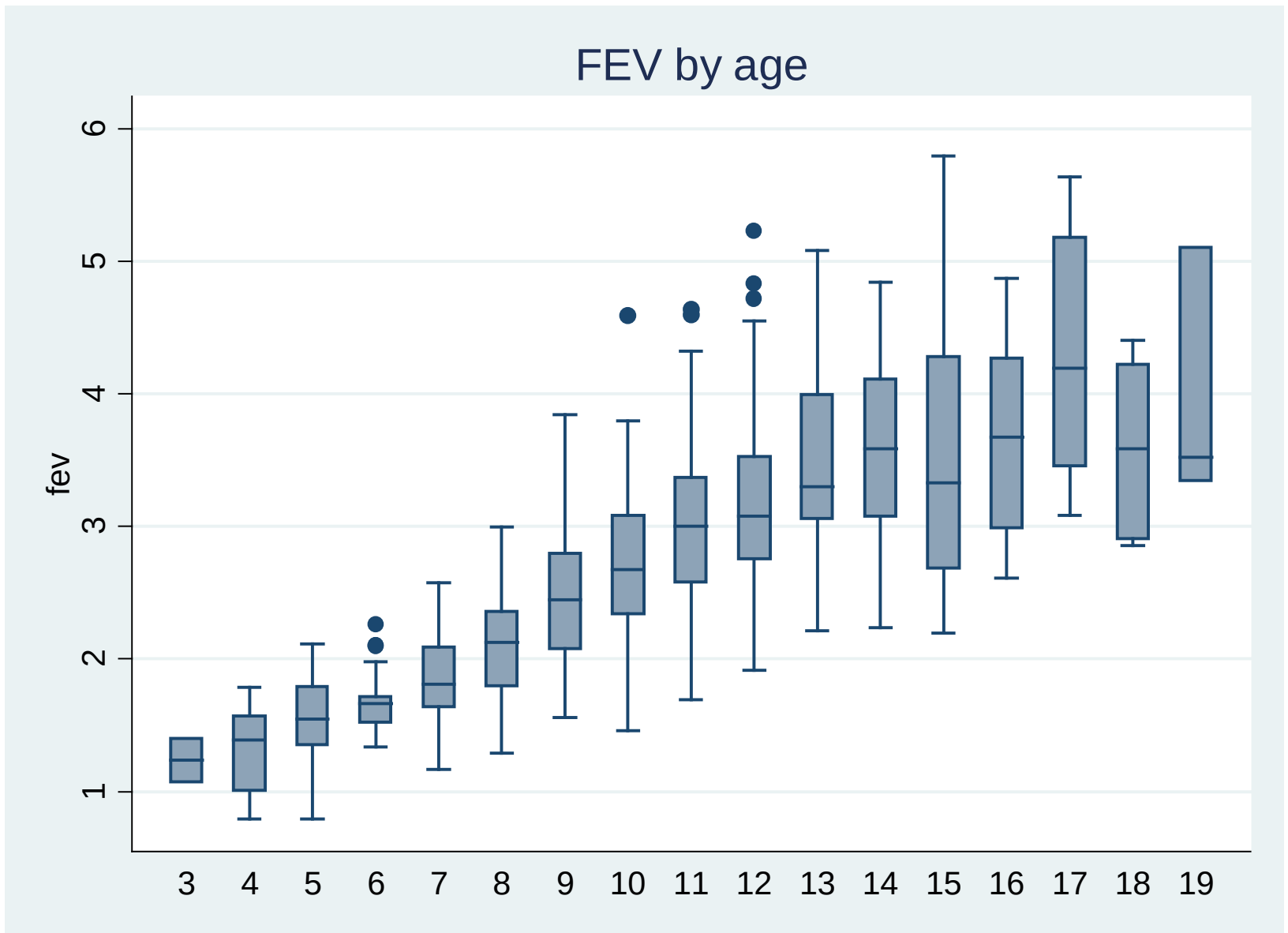
```
. spearman fev age
```

```
Number of obs =      654  
Spearman's rho =      0.7984
```

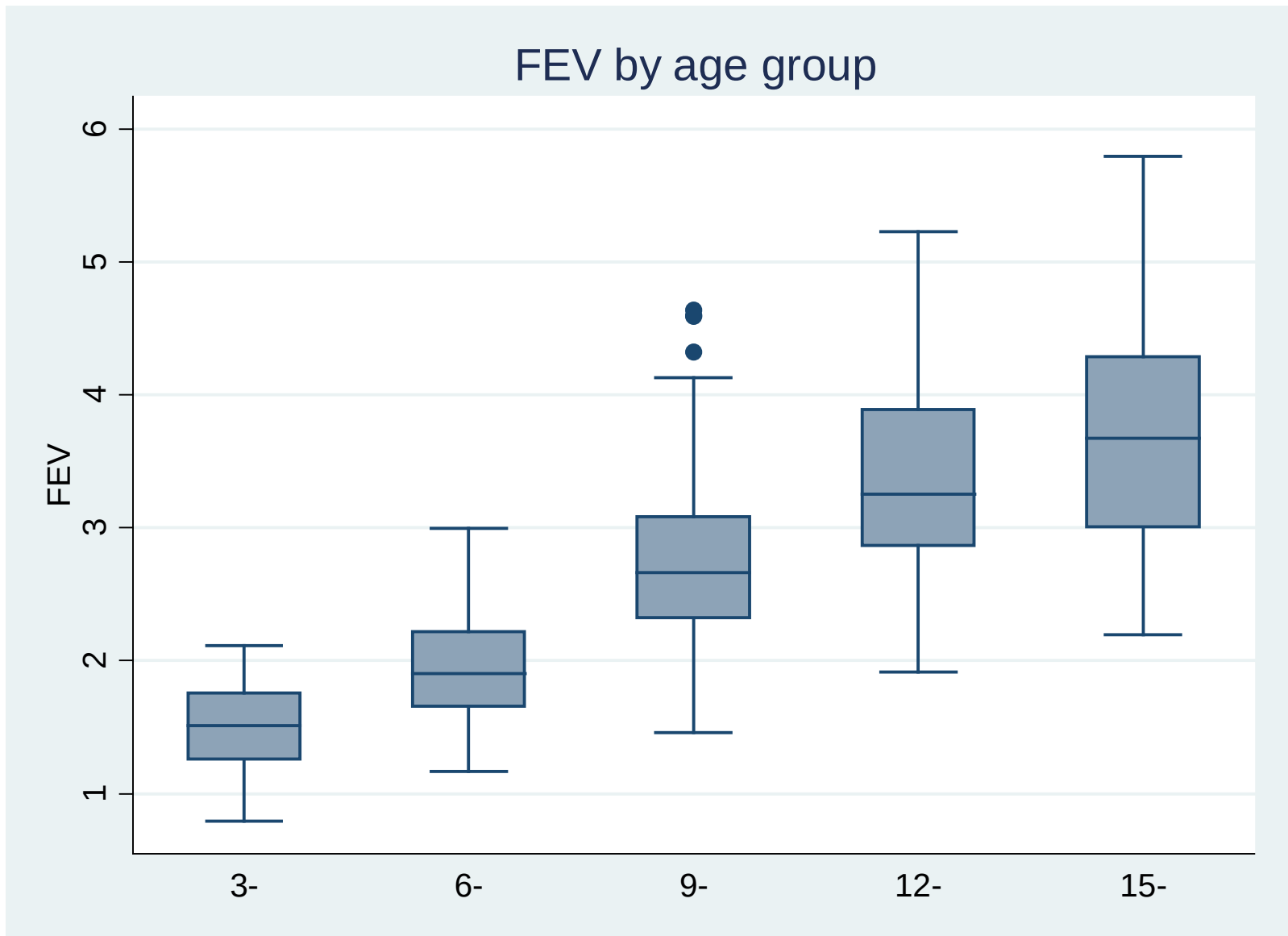
```
Test of Ho: fev and age are independent  
Prob > |t| =      0.0000
```

# Concept of $\mu_{y/x}$ and $\sigma_{y/x}$

- Consider variables X and Y that are thought to be related
- You want to know how a difference in X translates to a difference in Y
- Plot X versus Y, but instead of using all values of X, categorize X into several categories
- What you get would look like a boxplot of Y by the grouped values of X
- Each of the groups of X has a mean of Y  $\mu_{y/x}$  and a standard deviation  $\sigma_{y/x}$



graph box fev, over(age) title(FEV by age)



```
egen agecat = cut(age), at(0,3,6,9,12,15,20) label  
graph box fev, over(agecat) title(FEV by age group) ytitle(FEV)
```



```
. tabstat fev, by(agecat) s(n min median max mean sd)
```

Summary for variables: fev  
by categories of: agecat

| agecat | N   | min   | p50    | max   | mean     | sd       |
|--------|-----|-------|--------|-------|----------|----------|
| 3-     | 39  | .791  | 1.514  | 2.115 | 1.472385 | .3346982 |
| 6-     | 176 | 1.165 | 1.901  | 2.993 | 1.943727 | .3885005 |
| 9-     | 265 | 1.458 | 2.665  | 4.637 | 2.71723  | .5866867 |
| 12-    | 125 | 1.916 | 3.255  | 5.224 | 3.384576 | .7326963 |
| 15-    | 49  | 2.198 | 3.674  | 5.793 | 3.710143 | .8818795 |
| Total  | 654 | .791  | 2.5475 | 5.793 | 2.63678  | .8670591 |

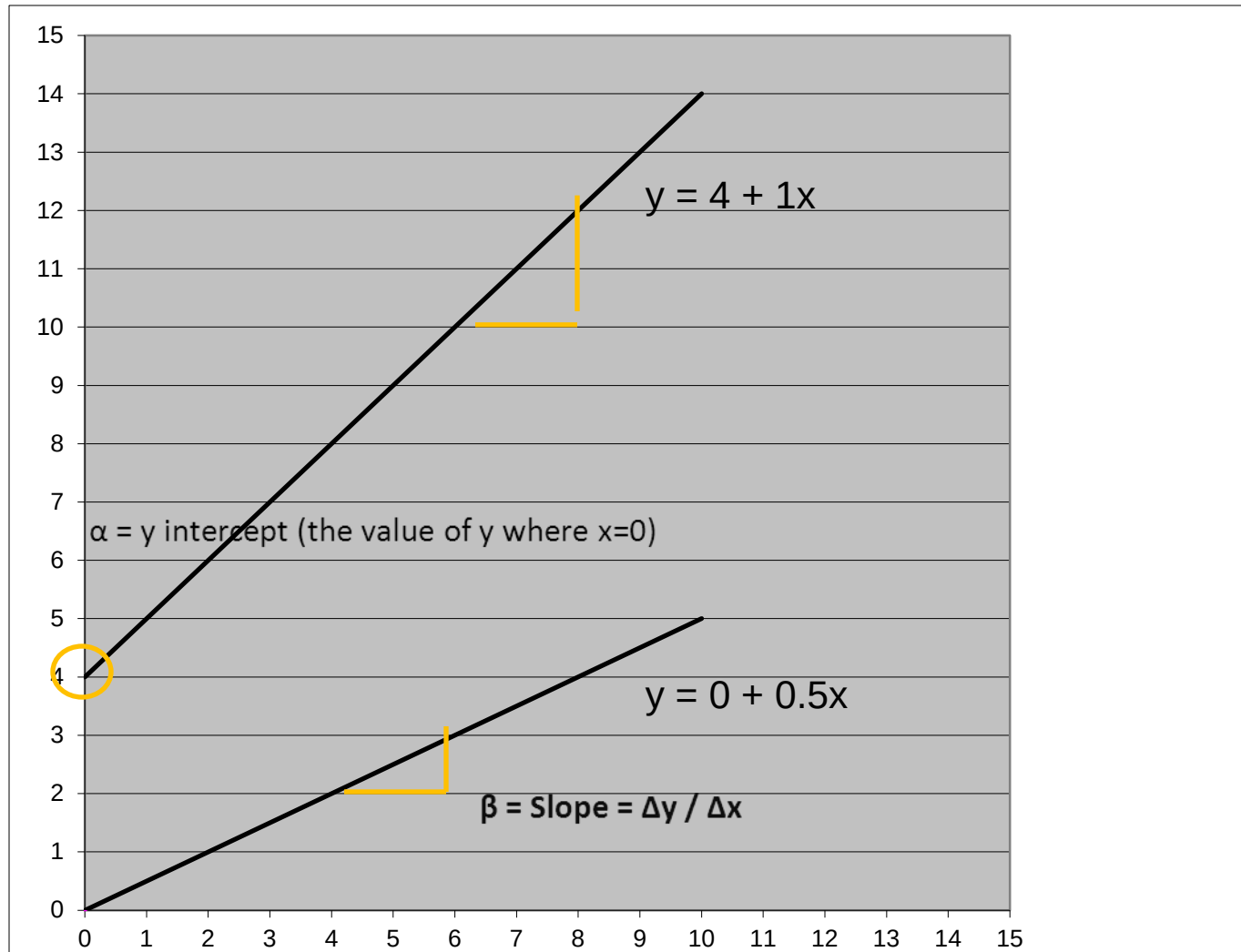


# Simple linear regression

- The method allows us to investigate the effect of a difference in the *explanatory* variable on the *response* variable.
- Equivalent terms:
  - Response variable, dependent variable, outcome variable, Y
  - Explanatory variable, independent variable, predictor variable, correlate, X
- Here it matters which variable is X and which variable is Y
- Y is the variable that you want to predict, or better understand with X (we regress Y on X)



# The equation of a straight line

$$y = \alpha + \beta x$$




# Simple linear regression

- Population regression equation is defined as

$$\mu_{y/x} = \alpha + \beta x$$

- This is the equation of a straight line
- $\alpha$  and  $\beta$  are constants and are called the coefficients of the equation

# Simple linear regression

- $\alpha$  is the y-intercept of the line  $y = \alpha + \beta x$
- Plugging back in to the population regression equation,

$\alpha$  is the mean value of Y when  $X=0$

$$\mu_{y|0} = \alpha + \beta X = \alpha$$

# Simple linear regression

- $\beta$  is the slope of the line
- $\beta$  is the change in the mean value of  $y$  that corresponds to a one-unit increase in  $X$

- E.g.  $X=3$  vs.  $X=2$

$$\mu_{y/3} - \mu_{y/2} = (\alpha + \beta*3) - (\alpha + \beta*2) = \beta$$

# Simple linear regression

- Even if there is a linear relationship between  $Y$  and  $X$  in theory, there will be some variability in the population
- At each value of  $X$ , there is a range of  $Y$  values, with a mean  $\mu_{y/x}$  and a standard deviation  $\sigma_{y/x}$
- We note this by including an error term,  $\varepsilon$ , in our regression equation



# Simple linear regression

- The linear regression equation is  $y = \alpha + \beta x + \varepsilon$
- The error,  $\varepsilon$ , is the distance a sample value  $y$  has from the population regression line

$$y = \alpha + \beta x + \varepsilon$$

$$\mu_{y/x} = \alpha + \beta x \text{ (population regression line)}$$

$$\text{so } y - \mu_{y/x} = \varepsilon$$

# Simple linear regression

- Assumptions of linear regression
  - X's are measured without error
    - Violations of this cause the coefficients to attenuate toward zero
  - For each value of X, the Y's are normally distributed with mean  $\mu_{y/x}$  and standard deviation  $\sigma_{y/x}$
  - The regression equation is correct  $\mu_{y/x} = \alpha + \beta x$

# Simple linear regression

- Assumptions of linear regression (continued)
  - Homoscedasticity – the standard deviation of  $y$  at each value of  $X$  is constant;  $\sigma_{y/x}$  the same for all values of  $X$ 
    - The opposite of homoscedasticity is heteroscedasticity
    - This is similar to the equal variance issue that we saw in ttests and ANOVA
  - All the  $y_i$  's are independent
    - You can't guess the  $y$  value for one person (or observation) based on the outcome of another
- Note that we do not need the  $X$ 's to be normally distributed, just the  $Y$ 's at each value of  $X$



# Least squares

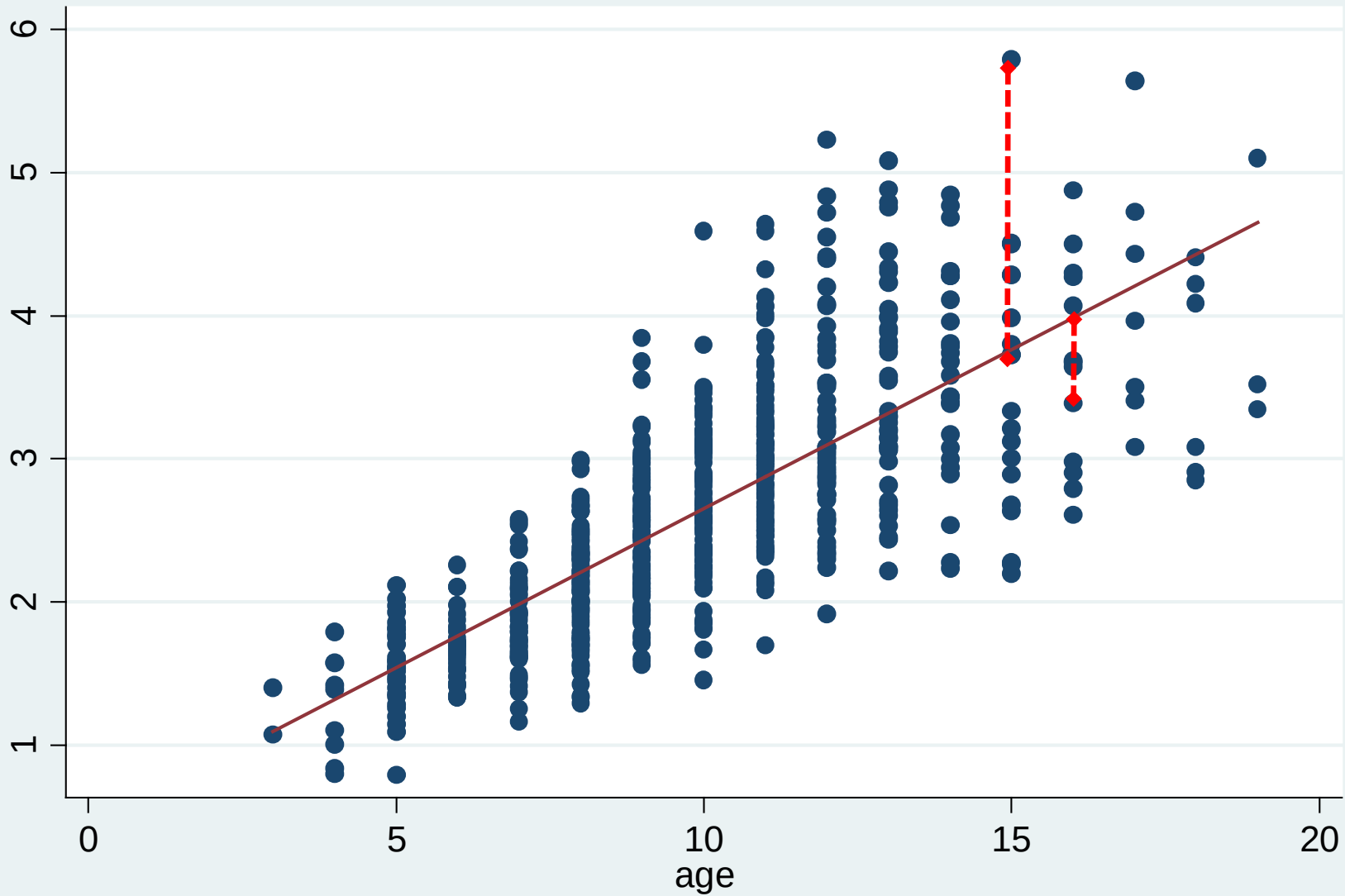
- We estimate the coefficients of the population regression line ( $\alpha$  and  $\beta$ ) using our sample of measurements of Y and X
- We have a set of data, where the points are  $(y_i, x_i)$ , and we want to put a line through them
- The distance from a data point  $(x_i, y_i)$  to the line at  $x_i$  is called the residual,  $e_i$

$$e_i = y_i - \hat{y}_i$$

$\hat{y}_i$  is y-value of the regression line at  $x_i$

It is the predicted value of y at each x

FEV versus age



# Simple linear regression

- The regression line equation is  $\hat{y} = \hat{\alpha} + \hat{\beta}x$
- The “best” line is the one that finds the  $\alpha$  and  $\beta$  that minimize the sum of the squared residuals  $\sum e_i^2$  (hence the name “least squares”)
- We are minimizing the sum of the squares of the residuals, called the error sum of squares or the residual sum of squares

$$\begin{aligned}\sum_{i=1}^n e_i^2 &= \sum_{i=1}^n (y_i - \hat{y}_i)^2 \\ &= \sum_{i=1}^n [y_i - (\hat{\alpha} + \hat{\beta}x_i)]^2\end{aligned}$$



# Simple linear regression

- The solution to this minimization is

$$\hat{\beta} = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sum_{i=1}^n (x_i - \bar{x})^2} \quad \text{and} \quad \hat{\alpha} = \bar{y} - \hat{\beta}\bar{x}$$

- These estimates are calculated directly from the x's and y's

# Simple linear regression example: Regression of FEV on age

$$\text{FEV} = \hat{\alpha} + \beta \text{ age}$$

regress yvar xvar

. regress fev age

| Source   | SS         | df  | MS         | Number of obs = 654    |  |  |
|----------|------------|-----|------------|------------------------|--|--|
| Model    | 280.919154 | 1   | 280.919154 | F( 1, 652) = 872.18    |  |  |
| Residual | 210.000679 | 652 | .322086931 | Prob > F = 0.0000      |  |  |
| Total    | 490.919833 | 653 | .751791475 | R-squared = 0.5722     |  |  |
|          |            |     |            | Adj R-squared = 0.5716 |  |  |
|          |            |     |            | Root MSE = .56753      |  |  |

| fev   | Coef.    | Std. Err. | t     | P> t  | [95% Conf. Interval] |          |
|-------|----------|-----------|-------|-------|----------------------|----------|
| age   | .222041  | .0075185  | 29.53 | 0.000 | .2072777             | .2368043 |
| _cons | .4316481 | .0778954  | 5.54  | 0.000 | .278692              | .5846042 |

$\beta$  = Coef for age

$\hat{\alpha}$  = \_cons (short for constant)

# Interpretation of the parameter estimates

- The least squares estimate is

$$\hat{y} = 0.432 + 0.222 x$$

- The intercept, 0.432 is the fitted value of y (FEV) for x (age) = 0
- The slope, 0.222 is the change in FEV corresponding to a change of 1 year in age. So a child with age=10 would have an FEV that is (on average) 0.222 higher than someone age 9. And the same for age 7 vs. 6, etc.

# Simple linear regression – hypothesis testing

- We want to know if there is a relationship between  $X$  and  $Y$ .
  - If there is no relationship then the value of  $Y$  does not change with the value of  $X$ , and  $\beta=0$ .
  - Therefore  $\beta=0$  is our null hypothesis.
- This is mathematically equivalent to the null hypothesis that the correlation  $\rho=0$ .
- We can also calculate a 95% confidence interval for  $\beta$

# Inference for regression coefficients

- We want to use the least squares regression line  $\hat{y} = \hat{\alpha} + \hat{\beta}x$  to make inference about the population regression line  $\mu_{y/x} = \alpha + \beta x$
- If we took repeated samples in which we measured  $x$  and  $y$  together and calculated the least squares estimates, we would have a distribution for the estimates  $\hat{\alpha}$  and  $\hat{\beta}$



# Inference for regression coefficients

- The standard error of the estimates are

$$se(\hat{\beta}) = \frac{\sigma_{y|x}}{\sqrt{\sum_{i=1}^n (x_i - \bar{x})^2}}$$

$$se(\hat{\alpha}) = \sigma_{y|x} \sqrt{\frac{1}{n} + \frac{\bar{x}^2}{\sum_{i=1}^n (x_i - \bar{x})^2}}$$

where we estimate  $\sigma_{y|x}$  with  $s_{y|x}$

$$\text{and } s_{y|x} = \sqrt{\frac{\sum_{i=1}^n (y_i - \hat{y}_i)^2}{n-2}}$$

# Inference for regression coefficients

- We can use these to test the null hypothesis  $H_0: \beta = \beta_0$  against the alternative  $H_0: \beta \neq \beta_0$
- The test statistic for this is  $t = \frac{\hat{\beta} - \beta_0}{\text{se}(\hat{\beta})}$
- And it follows the t distribution with  $n-2$  degrees of freedom under the null hypothesis

# Inference for regression coefficients

- When  $\beta_0=0$  , i.e. testing  $H_0: \beta =0$  , this is equivalent to testing  $\mu_{y/x} = \alpha + 0*x = \alpha$
- This is the same as testing the null hypothesis  $H_0: \rho=0$
- The regression slope and the correlation coefficient are related:  $\hat{\beta} = r \left( \frac{s_y}{s_x} \right)$
- 95% confidence intervals for  $\beta$   
(  $\beta - t_{n-2,.025} \text{se}(\beta)$  ,  $\beta + t_{n-2,.025} \text{se}(\beta)$  )

# Simple linear regression example: Regression of FEV on age

$$\text{FEV} = \hat{\alpha} + \beta \text{ age}$$

```
regress yvar xvar
```

```
. regress fev age
```

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|          |            |     |            | Adj R-squared = 0.5716 |  |  |
|          |            |     |            | Root MSE = .56753      |  |  |

| fev   | Coef.    | Std. Err. | t     | P> t  | [95% Conf. Interval] |          |
|-------|----------|-----------|-------|-------|----------------------|----------|
| age   | .222041  | .0075185  | 29.53 | 0.000 | .2072777             | .2368043 |
| _cons | .4316481 | .0778954  | 5.54  | 0.000 | .278692              | .5846042 |

# $R^2$

- A summary of the model fit is the coefficient of determination,  $R^2$
- $R^2 = r^2$ , i.e. the Pearson correlation coefficient squared
- $R^2$  ranges from 0 to 1, and measures the proportion of the variability in  $y$  that is explained by the regression of  $y$  on  $x$

# $R^2$

- $\sigma^2_{y|x} = (1 - \rho^2) \sigma^2_y$

If correlation=0, then  $\sigma^2_{y|x} = \sigma^2_y$

If correlation=1, then  $\sigma^2_{y|x} = 0$

- Substituting in sample values and rearranging:

$$R^2 = \frac{s_y^2 - s_{y|x}^2}{s_y^2}$$

- Looking at this formula illustrates how  $R^2$  represents the portion of the variability that is removed by performing the regression on X

# Simple linear regression: evaluating the model

$$\text{model sum of squares} = MSS = \sum_{i=1}^n (\hat{y}_i - \bar{y})^2$$

regress fev age

| Source   | SS         | df  | MS         |
|----------|------------|-----|------------|
| Model    | 280.919154 | 1   | 280.919154 |
| Residual | 210.000679 | 652 | .322086931 |
| Total    | 490.919833 | 653 | .751791475 |

Number of obs = 654

F( 1, 652) = 872.18

Prob > F = 0.0000

R-squared = 0.5722 ← = .7565<sup>2</sup>

Adj R-squared = 0.5716

Root MSE = .56753

| fev   | Coef.    | Std. Err. | t     | P> t  | [95% Conf. Interval] |          |
|-------|----------|-----------|-------|-------|----------------------|----------|
| age   | .222041  | .0075185  | 29.53 | 0.000 | .2072777             | .2368043 |
| _cons | .4316481 | .0778954  | 5.54  | 0.000 | .278692              | .5846042 |

$$\text{residual sum of squares} = RSS = \sum_{i=1}^n (y_i - \hat{y}_i)^2$$

$$\text{total sum of squares} = TSS = MSS + RSS$$

$$= \sum_{i=1}^n (y_i - \bar{y})^2$$

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

- Notation note:
  - Biostat 208 textbook Vittinghoff et al. use slightly different notation
  - The regression line notation we are using is

$$\hat{y} = \hat{\alpha} + \beta x$$

Vittinghoff et al. uses

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



# For next time

- Read Pagano and Gauvreau
  - Pagano and Gauvreau Chapter 17-18 (review)
  - Pagano and Gauvreau Chapter 18-19