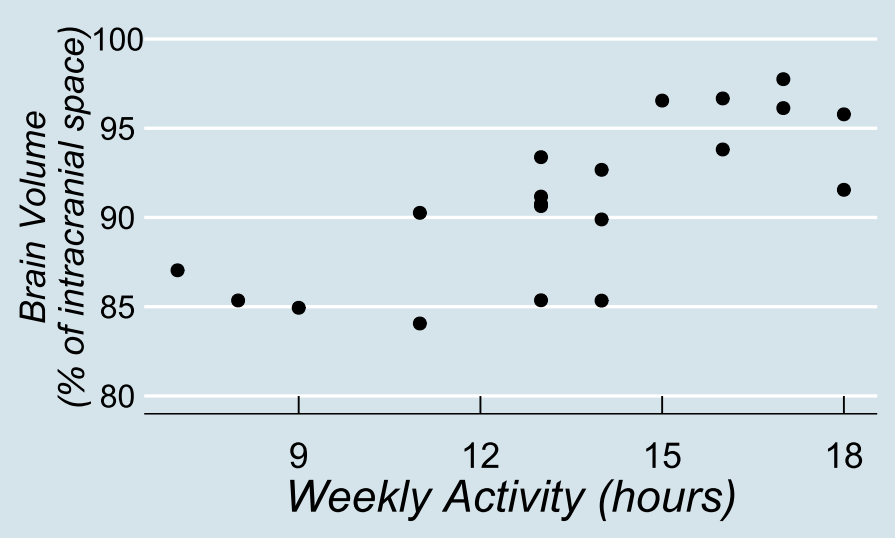






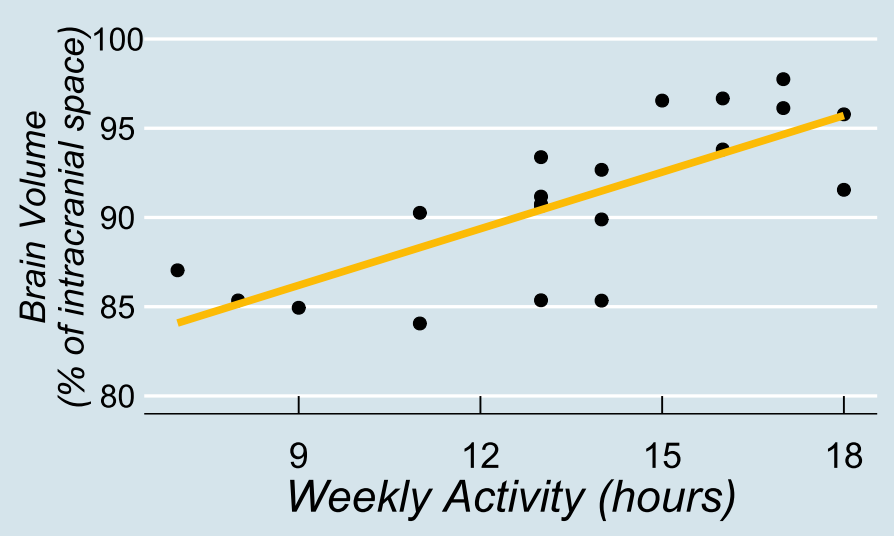
# Exercising our brains



$r = 0.7488, p = 0.0001$



# Exercising our brains (2)

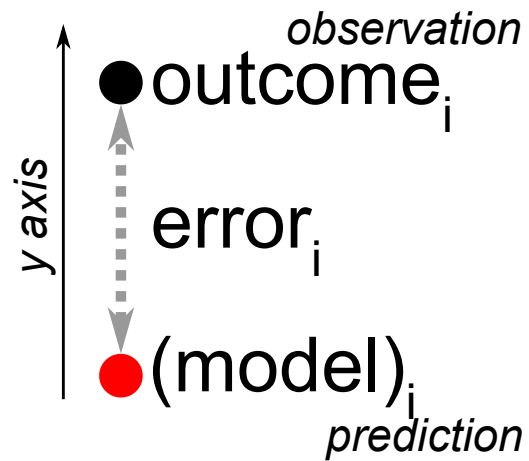


"for every extra 1 hour more weekly activity, brain volume increases by 1.06 (% of intracranial space)"



# The Only Equation You Will Ever Need

$$\text{outcome}_i = (\text{model})_i + \text{error}_i$$



# The Only Equation You Will Ever Need

$$\text{outcome}_i = (\text{model})_i + \text{error}_i$$

- to get any further, we need to make *assumptions*
- nature of the **model**
- nature of the **errors**

(linear)

(normal)

# A Linear Model

$$\text{outcome}_i = (\text{model})_i + \text{error}_i$$

$$y_i = b_0 \cdot 1 + b_1 \cdot x_i + \epsilon_i$$

so the linear model itself is...

$$\hat{y}_i = b_0 \cdot 1 + b_1 \cdot x_i$$

$$\mathbf{y} \sim \mathbf{1} + \mathbf{X}$$

# A Linear Model

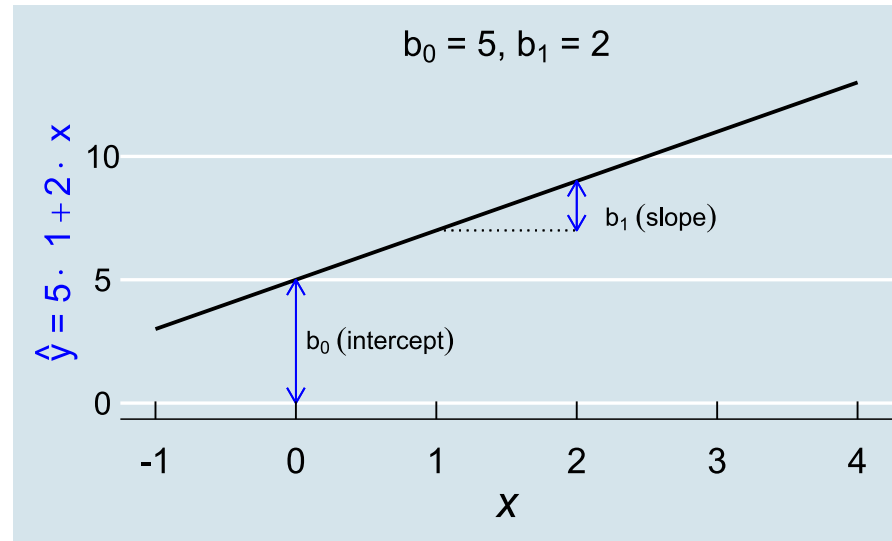
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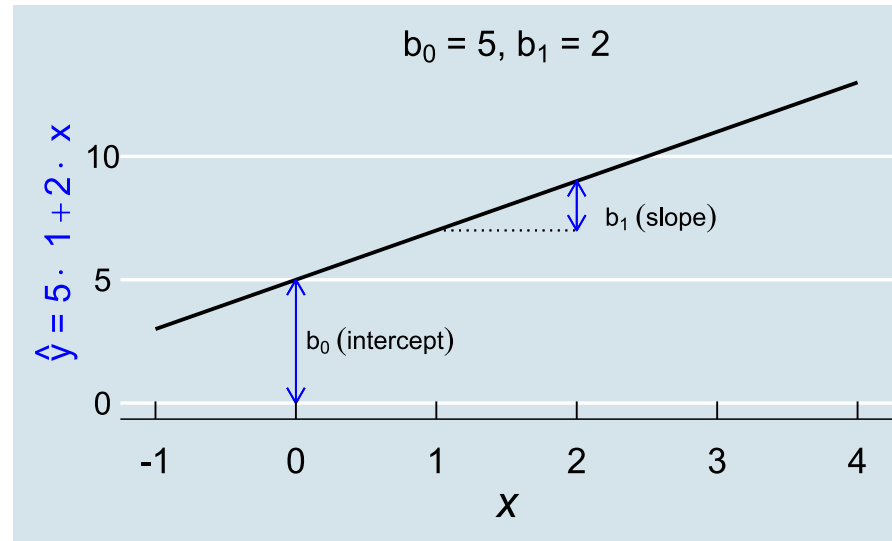
so the linear model itself is...

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$$y \sim 1 + x$$

$$\hat{y} = b_0 + b_1 \cdot x_i$$

$$y \sim X$$

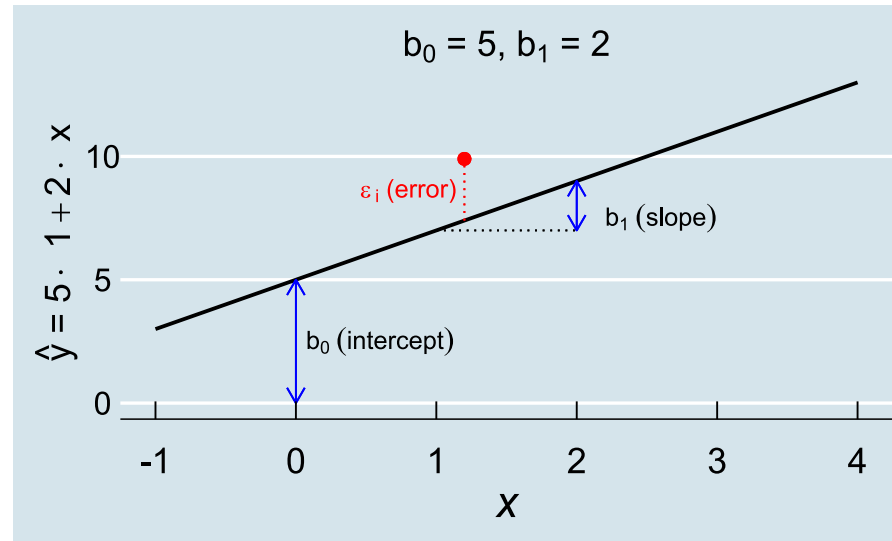


# Take An Observation

$$x_i = 1.2, y_i = 9.9$$

$$\hat{y}_i = b_0 + b_1 \cdot x_i = 7.4$$

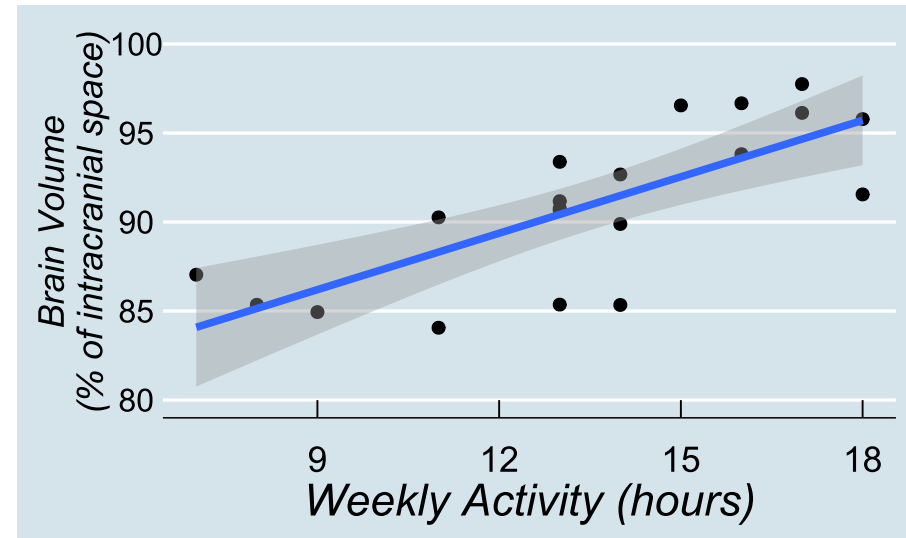
$$y_i = \hat{y}_i + \epsilon_i = 7.4 + 2.5$$



# More Brain Exercises

"for every extra 1 hour more weekly activity, brain volume increases by 1.06 (% of intracranial space)"

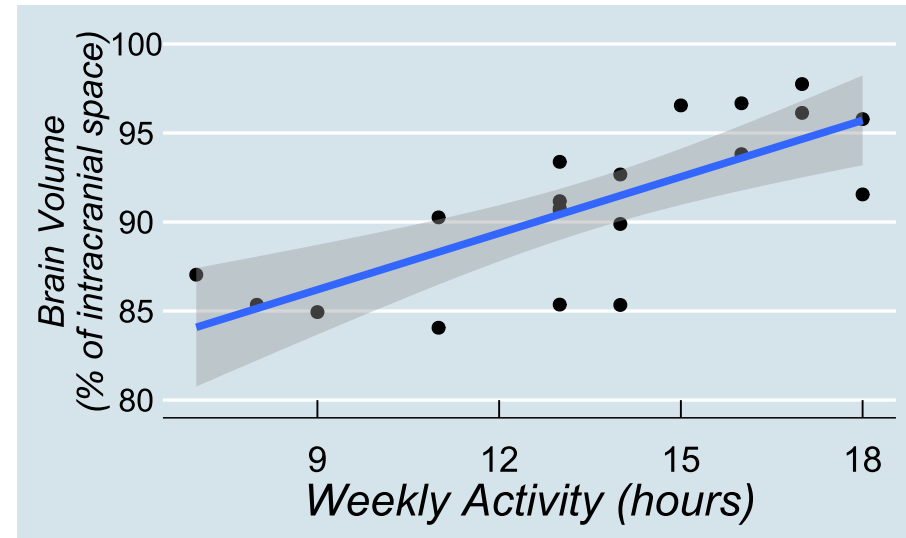
```
+ geom_smooth(method="lm")
```



# More Brain Exercises

"for every extra 1 hour more weekly activity, brain volume increases by 1.06 (% of intracranial space)"

```
+ geom_smooth(method="lm")
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but how can we evaluate our model?

# Linear Models in R

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mod <- lm(brain_vol ~ weekly_actv, data=dat)
```

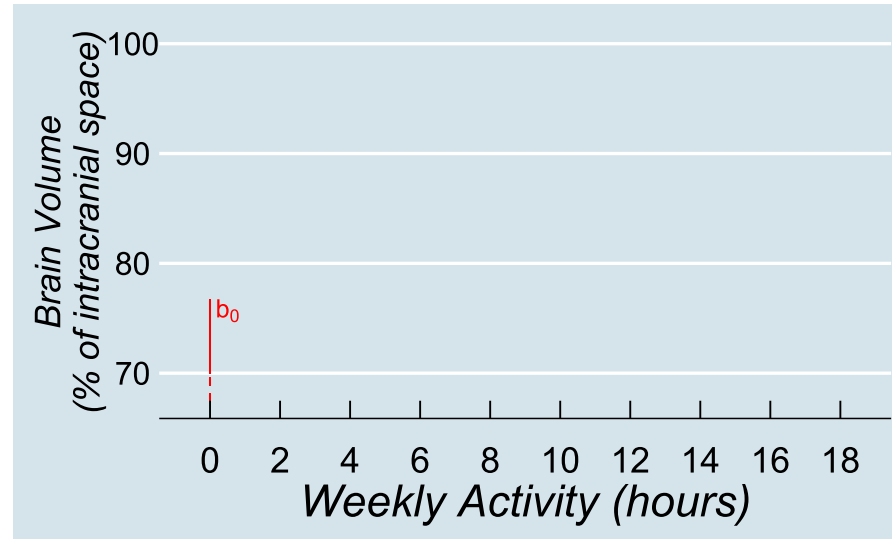
# Linear Models in R

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##
## Call:
## lm(formula = brain_vol ~ weekly_actv, data = dat)
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## Residuals:
##      Min       1Q   Median       3Q      Max
## -6.144 -1.342  0.274  2.199  4.009
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
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## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
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## Multiple R-squared:  0.561,    Adjusted R-squared:  0.536
## F-statistic:   23 on 1 and 18 DF,  p-value: 0.000145
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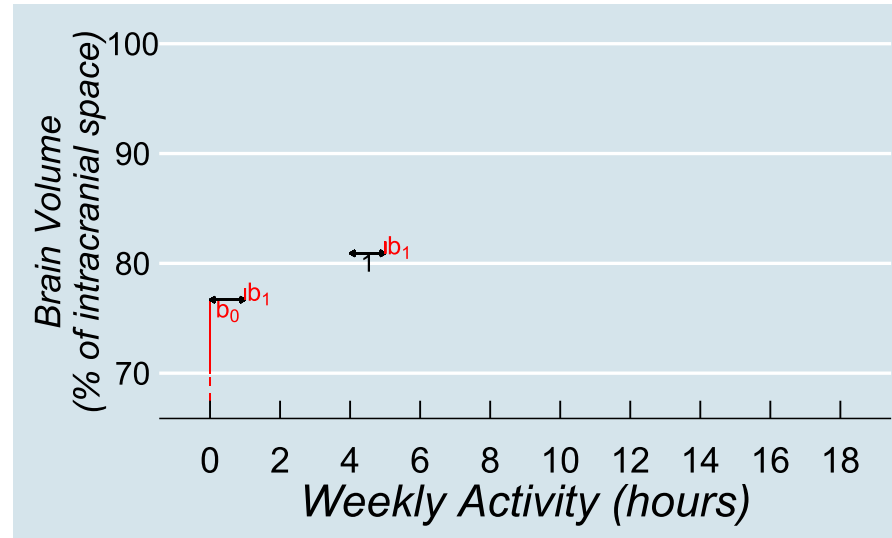
# Intercept and Slope Again

$$b_0 = 76.7; \quad b_1 = 1.06$$



# Intercept and Slope Again

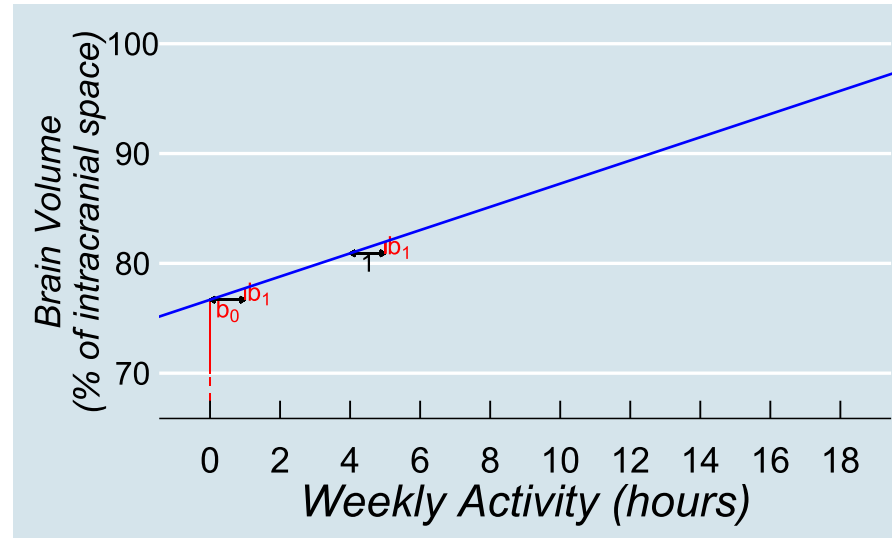
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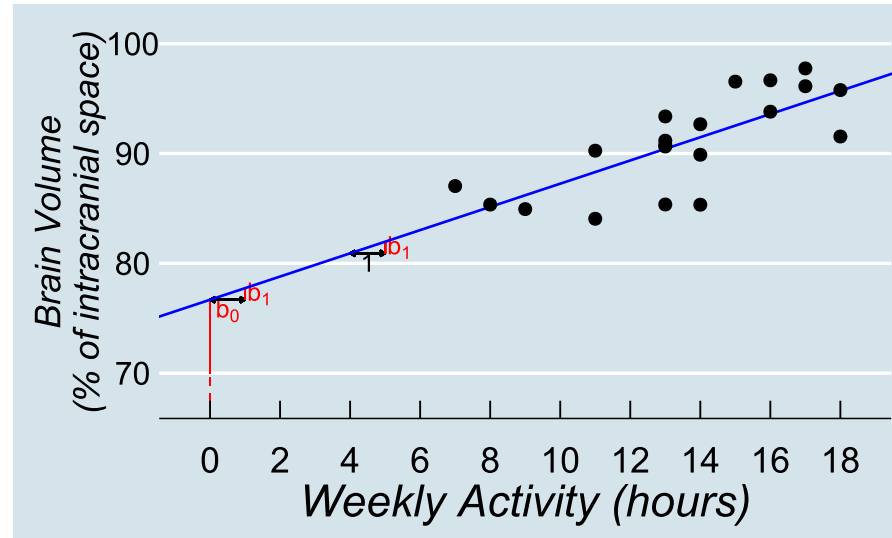
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# Are We Impressed?

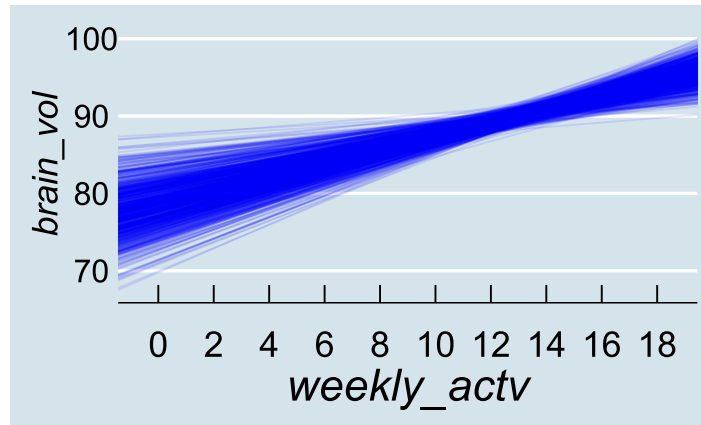
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- in NHST world, our pressing question is

# Are We Impressed?

- we have an intercept of 76.7 and a slope of 1.06
- in NHST world, our pressing question is

how likely would we have been to find these parameters under the null hypothesis?

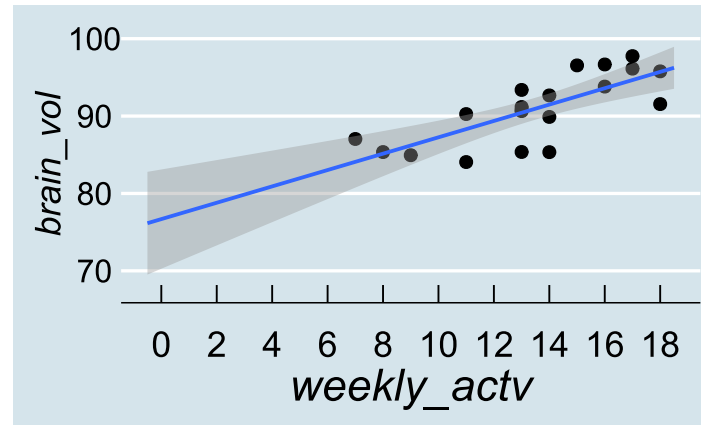
# Testing Chance



- repeatedly sampling 20~datapoints from the population
  - variability in *height* of line = variability in intercept ( $b_0$ )
  - variability in *angle* of line = variability in slope ( $b_1$ )



# We've Seen This Before



- shaded area represents "95% confidence interval"
  - if we repeatedly sampled 20 items from the population...
  - assumes that the 20 we have are the *best estimate* of the population

# The Good Old $t$ -Test

```
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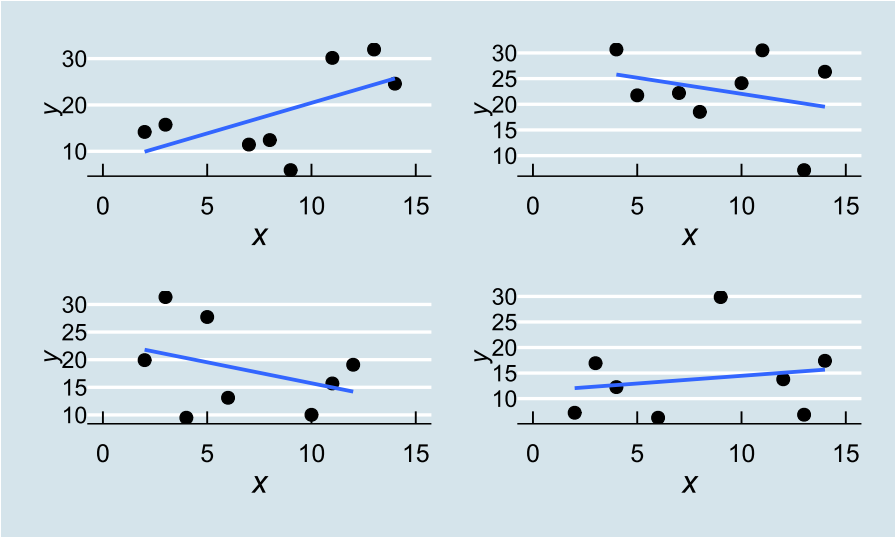
- for each model parameter we are interested in whether it is *different from zero*
- **intercept**: just like a mean
- **slope**: does the best-fit line differ from horizontal?

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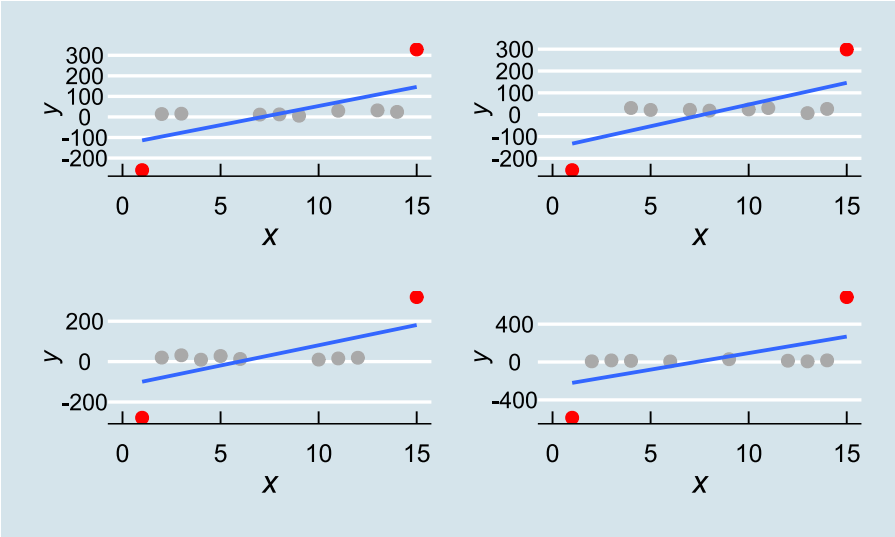
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- for each model parameter we are interested in whether it is *different from zero*
- **intercept**: just like a mean
- **slope**: does the best-fit line differ from horizontal?
- these are just (two-tailed) one-sample  $t$ -tests
  - **standard error** is the standard deviation of doing these lots of times
  - **t value** is  $\frac{\text{Estimate}}{\text{Std. Error}}$
  - to calculate  $p$ , we need to know the *degrees of freedom*

# Degrees of Freedom



# Degrees of Freedom



# Degrees of Freedom

- in fact we subtract 2 degrees of freedom because we "know" two things
  - intercept ( $b_0$ )
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# Degrees of Freedom

- in fact we subtract 2 degrees of freedom because we "know" two things
  - intercept ( $b_0$ )
  - slope ( $b_1$ )
- the remaining degrees of freedom are the *residual* degrees of freedom
- the *model* also has associated degrees of freedom
  - 2 (intercept, slope) - 1 (knowing one affects the other)

the models we have been looking at have 20 observations and 1 predictor

(1, 18) degrees of freedom

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```

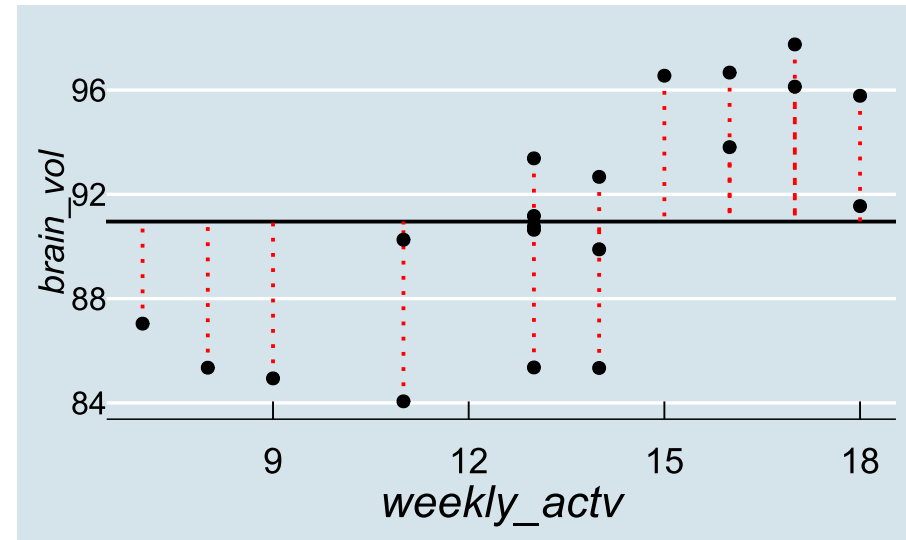
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```



# Total Sum of Squares

$$\text{total SS} = \sum (y - \bar{y})^2$$

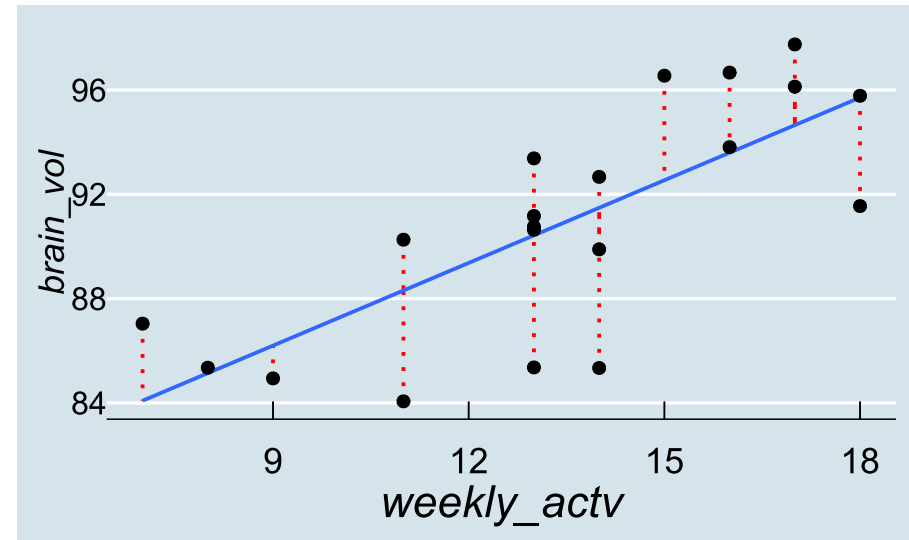
- sum of squares between observed  $y$  and mean  $\bar{y}$
- represents the total amount of variance in the model
- how much does the observed data vary from a model which says "there is no effect of  $x$ " (**null model**)?



# Residual Sum of Squares

$$\text{residual SS} = \sum (y - \hat{y})^2$$

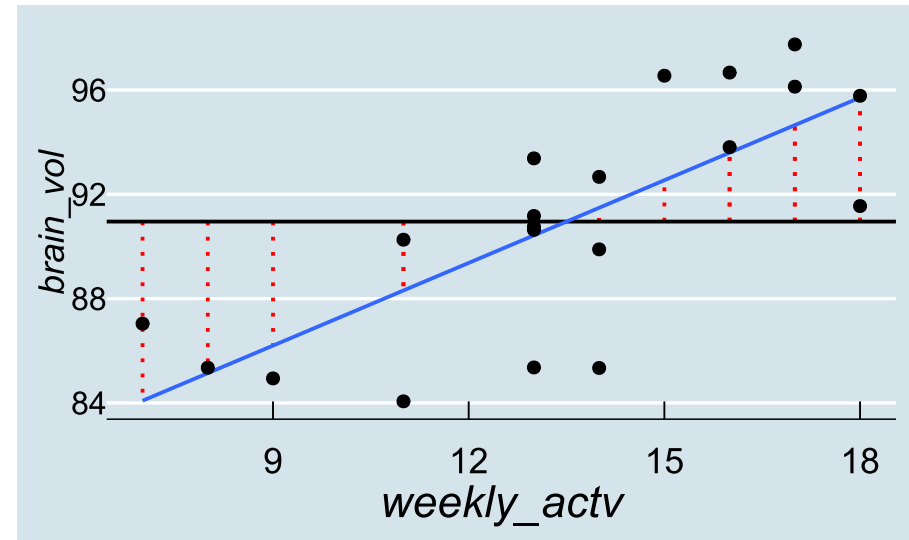
- sum of squared differences between observed  $y$  and predicted  $\hat{y}$
- represents the unexplained variance in the model
- how much does the observed data vary from the existing model?



# Model Sum of Squares

$$\text{model SS} = \sum (\hat{y} - \bar{y})^2$$

- sum of squared differences between predicted  $\hat{y}$  and mean  $\bar{y}$
- represents the additional variance explained by the current model over the null model

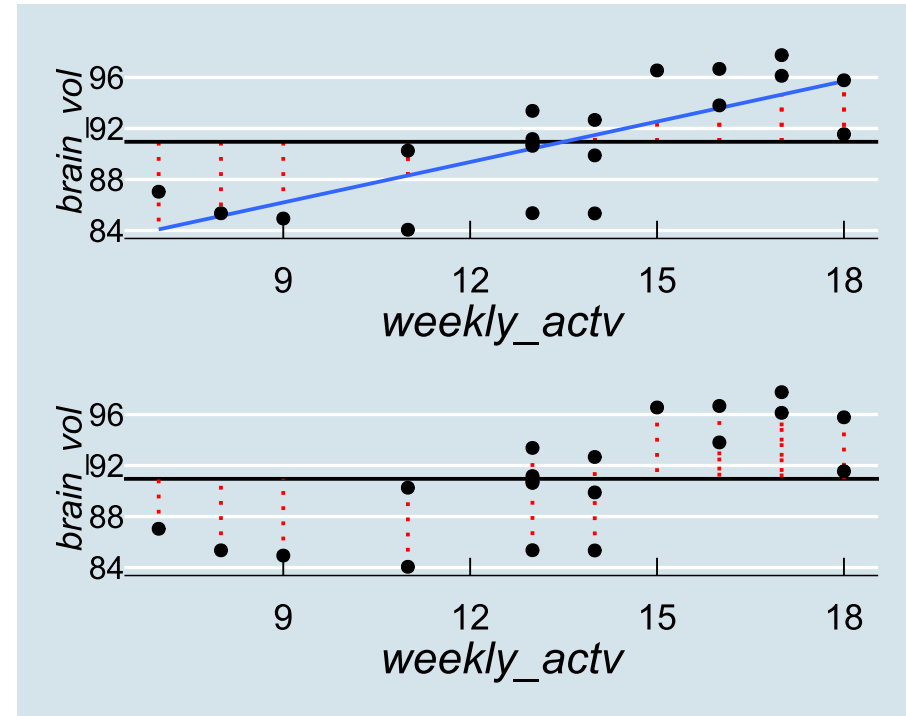


# Testing the Model: $R^2$

$$R^2 = \frac{\text{model SS}}{\text{total SS}} = \frac{\sum (\hat{y} - \bar{y})^2}{\sum (y - \bar{y})^2}$$

"how much the model improves over the null"

- $0 \leq R^2 \leq 1$
- we want  $R^2$  to be large
- for a single predictor,  $\sqrt{R^2} = |r|$



# Testing the Model: $F$

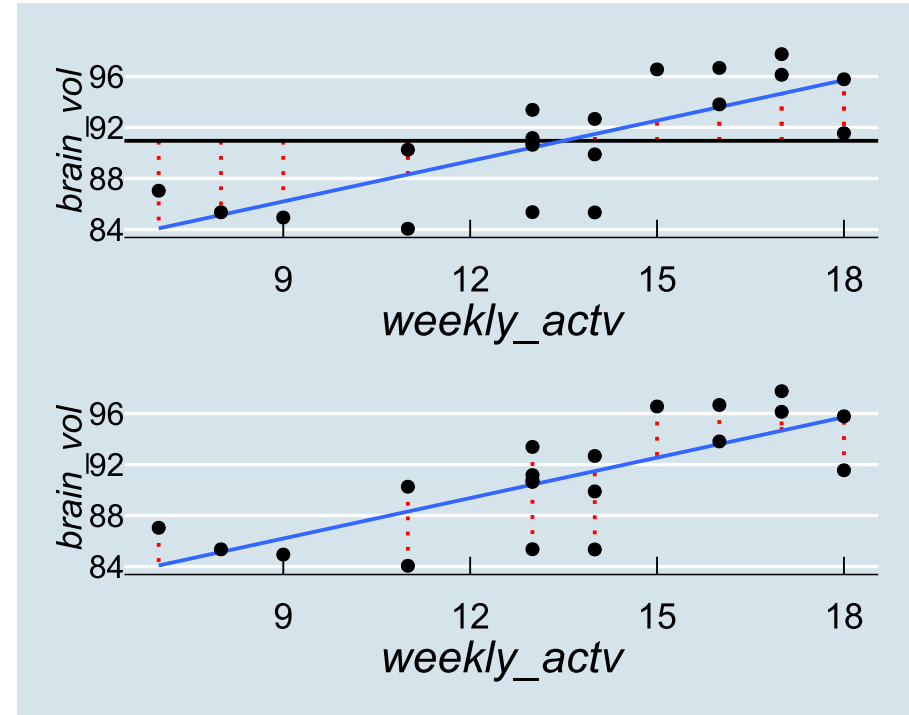
$F$  ratio depends on **mean squares**

$$F = \frac{\text{model MS}}{\text{residual MS}} = \frac{\sum (\hat{y} - \bar{y})^2 / \text{df}_m}{\sum (y - \hat{y})^2 / \text{df}_r}$$

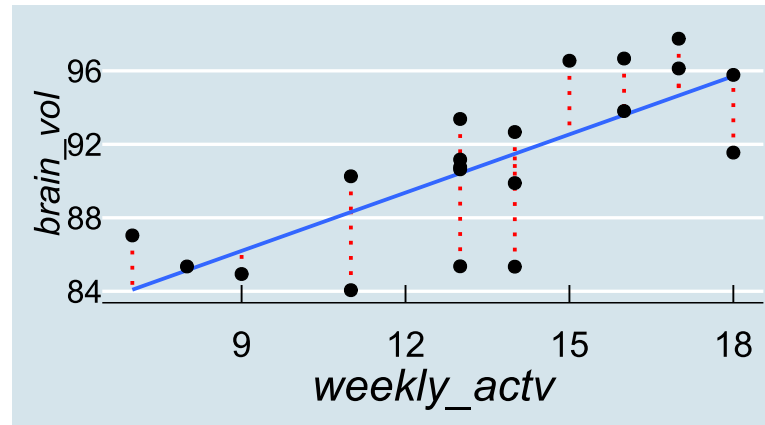
( $\text{MS}_x = \text{SS}_x / \text{df}_x$ )

"how much the model improves over chance"

- $0 < F$
- we want  $F$  to be large
- significance of  $F$  does not always equate to a large (or theoretically sensible) effect



# A Linear Model for 20 Brains



- a linear model describes the **best-fit line** through the data
- minimises the error terms  $\epsilon$  or **residuals**

# Two Types of Significance

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# The Good, the Bad, and the Ugly

- we can easily extend this approach
  - use more than one predictor
  - generalised linear model

- not a panacea
  - depends on *assumptions* about the data
  - depends on *decisions* about analysis



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  - depends on *assumptions* about the data
  - depends on *decisions* about analysis

- like other statistics, linear models don't tell you "about" your data
- they simply assess what is (un)likely to be due to chance
- the key to good statistics is *common sense and good interpretation*



