

# Week 4: Testing/Regression

Brandon Stewart<sup>1</sup>

Princeton

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<sup>1</sup>These slides are heavily influenced by Matt Blackwell, Adam Glynn and Jens Hainmueller.

# Where We've Been and Where We're Going...

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  - ▶ inference and estimator properties
  - ▶ point estimates, confidence intervals

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- Long Run
  - ▶ probability  $\rightarrow$  inference  $\rightarrow$  regression

Questions?

- 1 Testing: Making Decisions
  - Hypothesis testing
  - Forming rejection regions
  - P-values
- 2 Review: Steps of Hypothesis Testing
- 3 The Significance of Significance
- 4 Preview: What is Regression
- 5 Fun With Salmon
- 6 Nonparametric Regression
  - Discrete  $X$
  - Continuous  $X$
  - Bias-Variance Tradeoff
- 7 Linear Regression
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# Example

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- The trial included 345 patients with initial systolic blood pressure between 140-159.
- Each subject was assigned to take the drug combination for 16 weeks.
- Systolic blood pressure was measured on each subject before and after the treatment period.

# Example

Subject	SBP <sub>before</sub>	SBP <sub>after</sub>	Decrease
1			
2			
3			
4			
⋮			
345			

## Example

Subject	SBP <sub>before</sub>	SBP <sub>after</sub>	Decrease
1	147		
2	153		
3	142		
4	141		
⋮	⋮		
345	155		



## Example

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2	153	122	
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4	141	134	
⋮	⋮	⋮	
345	155	115	

## Example

Subject	SBP <sub>before</sub>	SBP <sub>after</sub>	Decrease
1	147	135	12
2	153	122	31
3	142	119	23
4	141	134	7
⋮	⋮	⋮	⋮
345	155	115	40

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Question: Should the FDA allow the drug to proceed to the next stage of testing?

# The FDA's Decision

We can think of the FDA's problem in terms of two dimensions:

- The true state of the world
- The decision made by the FDA

	Drug works	Drug doesn't work
FDA approves		
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- **Alternative Hypothesis:** Claim to be tested (research hypothesis)

Example: The drug does reduce blood pressure on average  
( $\mu_{decrease} > 0$ )

# More Examples

Null Hypothesis Examples ( $H_0$ ):

Alternative Hypothesis Examples ( $H_a$ ):

## More Examples

### Null Hypothesis Examples ( $H_0$ ):

- The drug does not change blood pressure on average ( $\mu_{decrease} = 0$ )

### Alternative Hypothesis Examples ( $H_a$ ):

- The drug does change blood pressure on average ( $\mu_{decrease} \neq 0$ )



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Back to the two dimensions of the FDA's problem:

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FDA approves (reject $H_0$ )	Correct	Type I error
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FDA approves (reject $H_0$ )	Correct	Type I error
FDA doesn't approve (don't reject $H_0$ )	Type II error	Correct

# Test Statistics, Null Distributions, and Rejection Regions

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**Null Distribution:** the sampling distribution of the statistic/test statistic assuming that the null is true.

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If we assume that the null hypothesis is true such that  $\mu = \mu_0$ , then

$$\begin{aligned}\bar{X} &\sim_{\text{approx}} N(\mu_0, S^2/n) \\ \frac{\bar{X} - \mu_0}{\frac{S}{\sqrt{n}}} &\sim_{\text{approx}} N(0, 1)\end{aligned}$$

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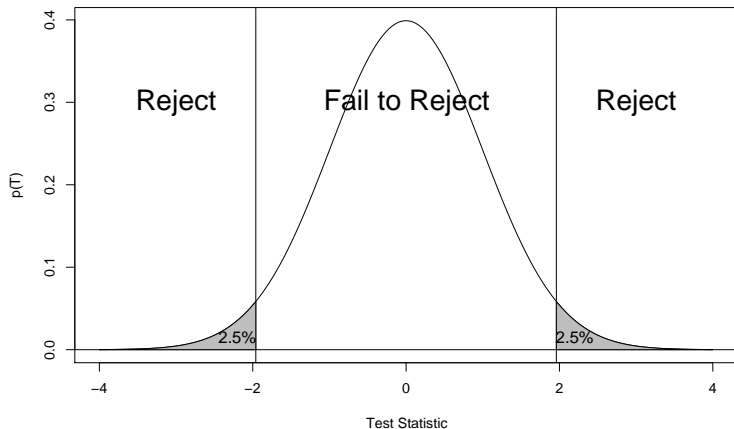
We usually pick an  $\alpha$  that we are comfortable with in advance, and using the null distribution for the test statistic and the alternative hypothesis, we define a **rejection region**.

Example: Suppose  $\alpha = 5\%$ , the test statistic is  $\frac{\bar{X} - \mu_0}{\frac{s}{\sqrt{n}}}$ , the null hypothesis is  $H_0 : \mu = \mu_0$ , and the alternative hypothesis is  $H_a : \mu \neq \mu_0$ .

# Two-sided rejection region

## Two-sided rejection region

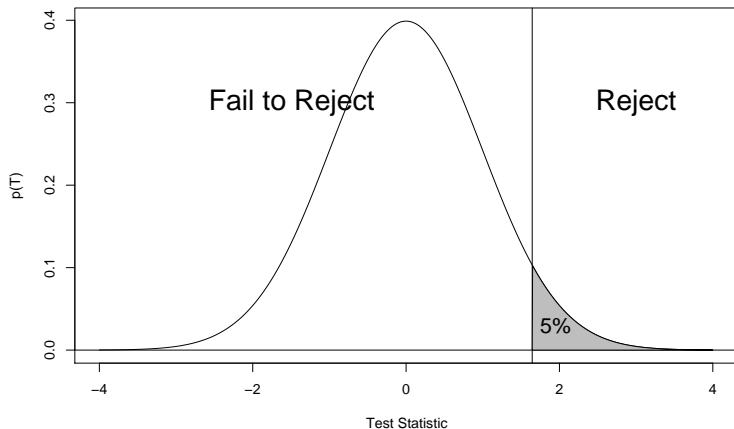
Rejection region with  $\alpha = .05$ ,  $H_0 : \mu = 0$ ,  $H_A : \mu \neq 0$ :



# One-sided Rejection Region

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Rejection region with  $\alpha = .05$ ,  $H_0 : \mu \leq 0$ ,  $H_A : \mu > 0$ :



## Example

So, should the FDA approve further trials?

Recall the null and alternative hypotheses:

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- $\bar{x} = 21.0$
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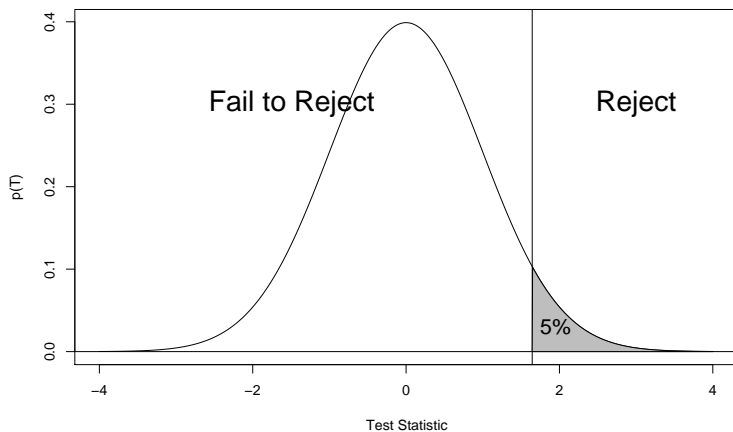
- $\bar{x} = 21.0$
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Therefore,

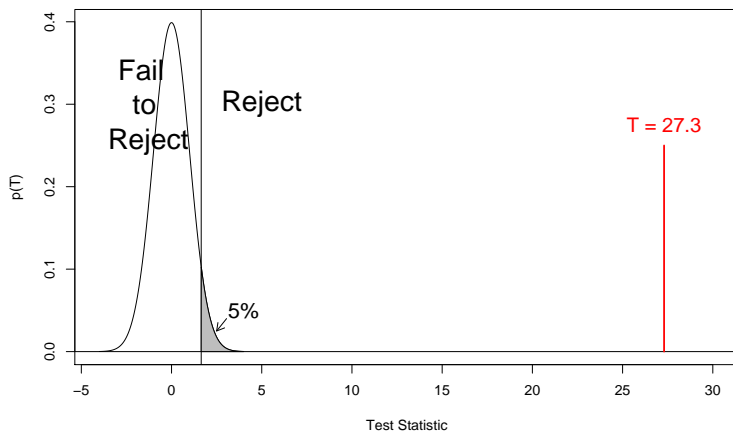
$$T = \frac{21.0 - 0}{\frac{14.3}{\sqrt{345}}} = 27.3$$

What is the decision?

## Rejection Region with $\alpha = .05$



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**P-value:** Assuming that the null hypothesis is true, the probability of getting something at least as extreme as our observed test statistic, where extreme is defined in terms of the alternative hypothesis.

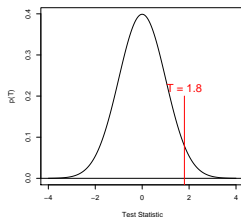
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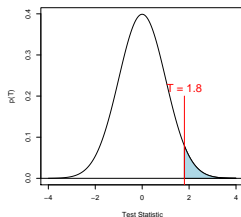
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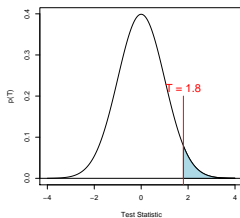
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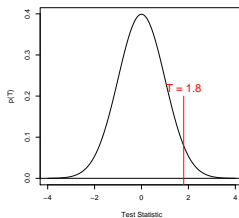
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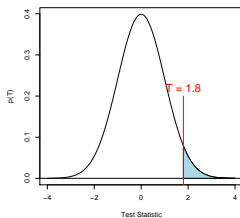
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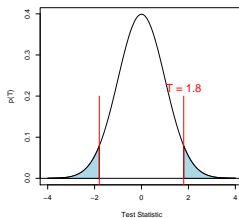
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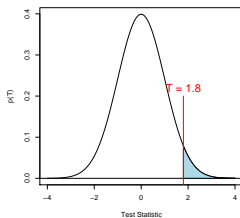


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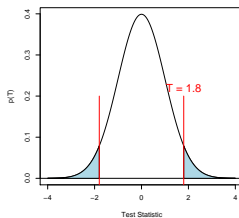
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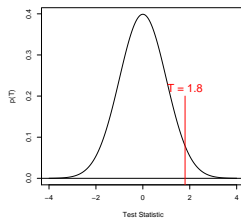
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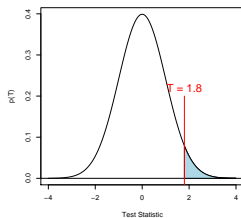
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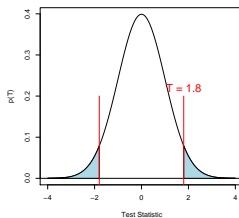
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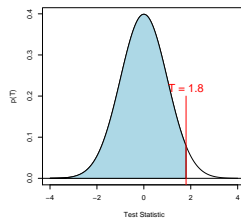
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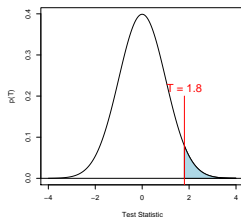
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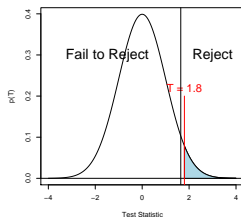
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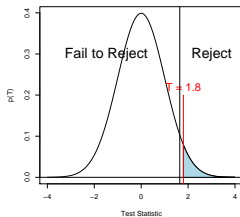
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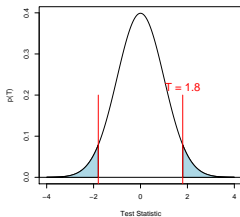
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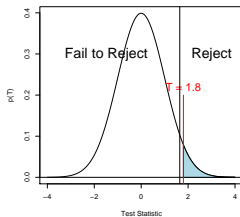




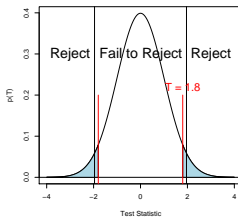
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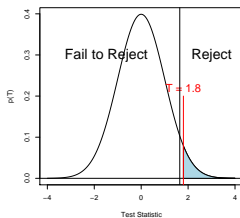
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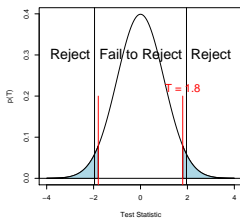
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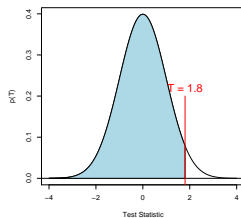
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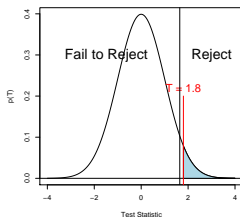
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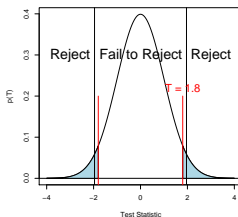
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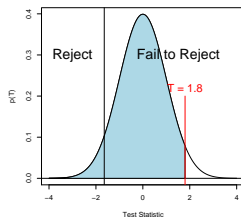
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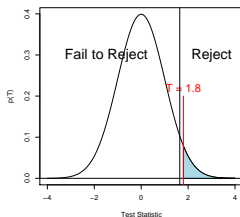
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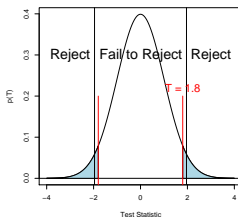
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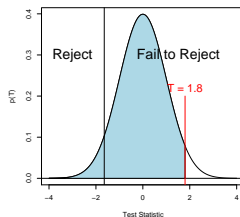
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If  $p < \alpha$ , then the test statistic falls in the rejection region for the  $\alpha$ -level test.

## Example 1

Recall the drug testing example, where  $H_0 : \mu_0 \leq 0$  and  $H_a : \mu_0 > 0$ :

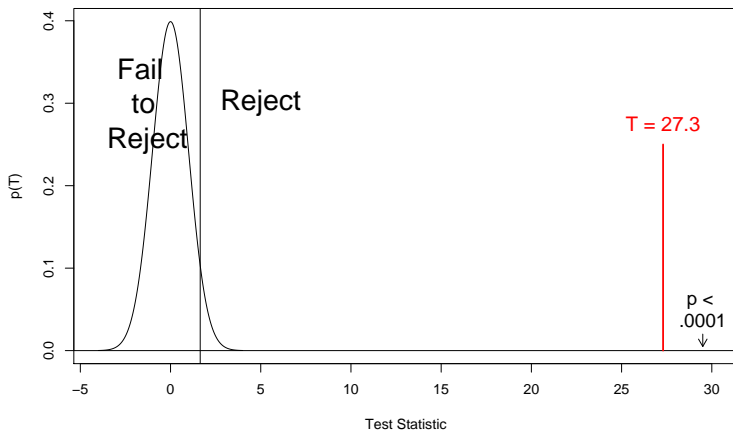
- $\bar{x} = 21.0$
- $s = 14.3$
- $n = 345$

Therefore,

$$T = \frac{21.0 - 0}{\frac{14.3}{\sqrt{345}}} = 27.3$$

What is the probability of observing a test statistic greater than 27.3 if the null is true?

# Example 1



## $\alpha$ Rejection Regions and $1 - \alpha$ CIs

Up to this point, we have defined rejection regions in terms of the test statistic.

In some cases, we can define an equivalent rejection region in terms of the parameter of interest.

For a two-sided, large-sample test, we reject if:

$$\frac{\bar{X} - \mu_0}{\frac{s}{\sqrt{n}}} > z_{\alpha/2} \text{ or } \frac{\bar{X} - \mu_0}{\frac{s}{\sqrt{n}}} < -z_{\alpha/2}$$

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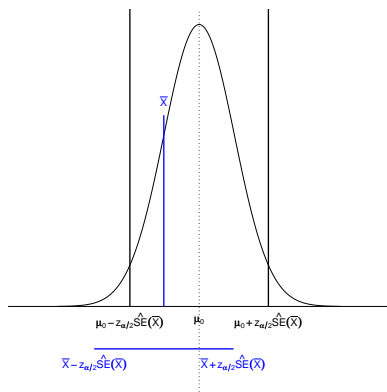
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Therefore, we can use the  $1 - \alpha$  CI to test the null hypothesis at the  $\alpha$  level.



## Another interpretation of CIs

The form of the “fail to reject” region of an  $\alpha$ -level hypothesis test is:

$$\left( \mu_0 - z_{\alpha/2} \times \frac{s}{\sqrt{n}}, \mu_0 + z_{\alpha/2} \times \frac{s}{\sqrt{n}} \right)$$

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So the  $1 - \alpha$  CI is the set of null hypotheses  $\mu_0$  that would not be rejected at the  $\alpha$  level.



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However, creating a decision rule which minimizes both types of errors at the same time is impossible. We therefore need to balance them.

## Hypothesis Testing: Error Types

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**Hypothesis testing** follows an analogous logic, where we want to decide whether to **reject** (= convict) or **fail to reject** (= acquit) a **null hypothesis** (= defendant) using sample data.

# Hypothesis Testing: Steps

		<i>Null Hypothesis (<math>H_0</math>)</i>	
		False	True
<i>Decision</i>	Reject	$1 - \beta$	$\alpha$
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- 4 Assuming  $H_0$  is true, derive the **null distribution** of  $T$  (e.g. standard normal)

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- 5 Using the **critical values** from a statistical table, evaluate how unusual the observed value of  $T$  is under the null hypothesis:
- ▶ If the probability of drawing a  $T$  **at least as extreme** as the observed  $T$  is less than  $\alpha$ , we reject  $H_0$ .  
(e.g. there are too many testimonies against the defendant for her to be innocent, so reject the hypothesis that she was innocent.)
  - ▶ Otherwise, we fail to reject  $H_0$ .  
(e.g. there is not enough evidence against the defendant, so give her the benefit of the doubt.)

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- We need to be careful to distinguish:
  - ▶ **practical significance** (e.g. a big effect)
  - ▶ **statistical significance** (i.e. we reject the null)
- In large samples even tiny effects will be significant, but the **results may not be very important substantively**. Always discuss both!

# Star Chasing (aka there is an XKCD for everything)

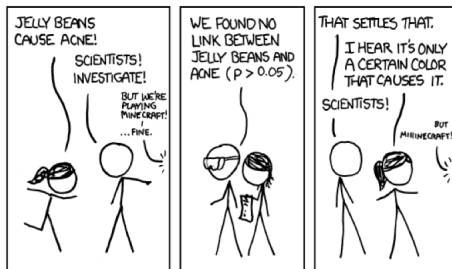
ARCHIVE  
FORUMS  
NEWS/BLAG  
STORE  
ABOUT

**xkcd** A WEBCOMIC OF ROMANCE,  
SARCASM, MATH, AND LANGUAGE.

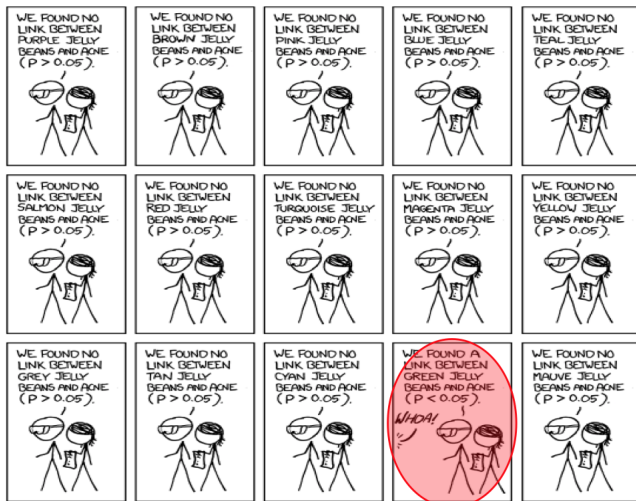
XKCD UPDATES EVERY MONDAY, WEDNESDAY, AND FRIDAY.

## SIGNIFICANT

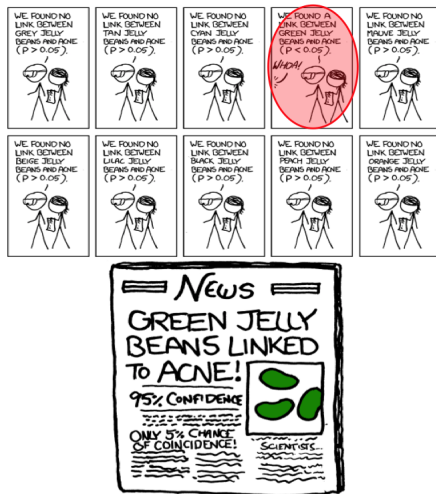
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- Illustration: randomly draw 21 variables, and run a regression of the first variable on the rest.

# Multiple Testing

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- Illustration: randomly draw 21 variables, and run a regression of the first variable on the rest.
- By design, no effect of any variable on any other, but when we run the regression:

# Multiple Test Example

```
##
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept) -0.0280393  0.1138198  -0.246  0.80605
## X2          -0.1503904  0.1121808  -1.341  0.18389
## X3           0.0791578  0.0950278   0.833  0.40736
## X4          -0.0717419  0.1045788  -0.686  0.49472
## X5           0.1720783  0.1140017   1.509  0.13518
## X6           0.0808522  0.1083414   0.746  0.45772
## X7           0.1029129  0.1141562   0.902  0.37006
## X8          -0.3210531  0.1206727  -2.661  0.00945 **
## X9          -0.0531223  0.1079834  -0.492  0.62412
## X10          0.1801045  0.1264427   1.424  0.15827
## X11          0.1663864  0.1109471   1.500  0.13768
## X12          0.0080111  0.1037663   0.077  0.93866
## X13          0.0002117  0.1037845   0.002  0.99838
## X14          -0.0659690  0.1122145  -0.588  0.55829
## X15          -0.1296539  0.1115753  -1.162  0.24872
## X16          -0.0544456  0.1251395  -0.435  0.66469
## X17           0.0043351  0.1120122   0.039  0.96923
## X18          -0.0807963  0.1098525  -0.735  0.46421
## X19          -0.0858057  0.1185529  -0.724  0.47134
## X20          -0.1860057  0.1045602  -1.779  0.07910 .
## X21           0.0021111  0.1081179   0.020  0.98447
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.9992 on 79 degrees of freedom
## Multiple R-squared:  0.2009, Adjusted R-squared:  -0.00142
## F-statistic: 0.993 on 20 and 79 DF,  p-value: 0.4797
```

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- But this is exactly what we expect:  $1/20 = 0.05$  of the tests are false positives at the 0.05 level
- Also note that  $2/20 = 0.1$  are significant at the 0.1 level. Totally expected!

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- The most prominent adjustments include:
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- It remains a heated debate.

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- Reporting p-values allows the researcher to separate the analysis from the decision.
- There is a close relationship between the results of an  $\alpha$  level hypothesis test and the coverage of a  $(1 - \alpha)\%$  confidence interval.

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- From here on out, we'll be interested in the relationships between variables. How does one variable change as we change the values of another variable? This question will be the bread and butter of the class moving forward.

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  - ▶ Social pressure mailer versus Civic Duty Mailer
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- Generally our goal is to understand how  $Y$  varies as a function of  $X$ :

$$Y = f(X) + \text{error}$$



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- Joint densities, covariance, and correlation were all ways to summarize the relationship between two variables.
- But these were population quantities and we only have samples, so we may want to estimate these quantities using their sample analogs (plug-in principle or analogy principle)

# Scatterplots

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- Shows graphically how two variables are related

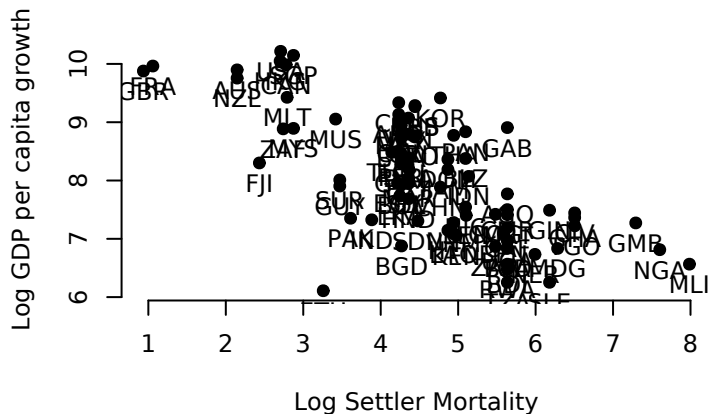


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Data from Acemoglu, Johnson and Robinson

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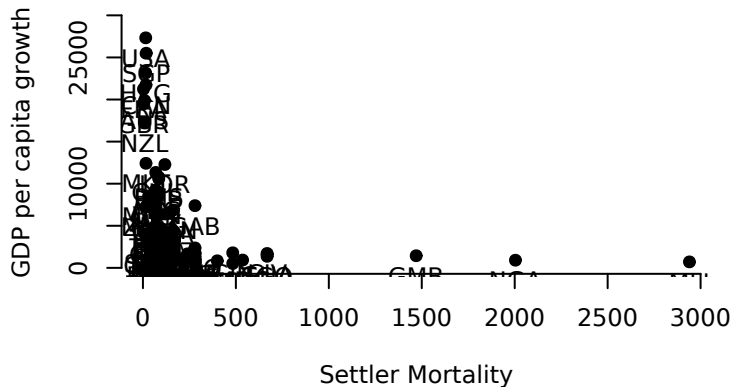
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## Definition (Sample Covariance)

The **sample covariance** between  $Y_i$  and  $X_i$  is

$$S_{XY} = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X}_n)(Y_i - \bar{Y}_n)$$

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$$\hat{\rho} = r = \frac{S_{XY}}{S_X S_Y} = \frac{\sum_{i=1}^n (X_i - \bar{X}_n)(Y_i - \bar{Y}_n)}{\sqrt{\sum_{i=1}^n (X_i - \bar{X}_n)^2 \sum_{i=1}^n (Y_i - \bar{Y}_n)^2}}$$

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- Once we have estimated  $E[Y|X]$ , we can use it for **prediction** and/or **causal inference**, depending on what assumptions we are willing to make

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- Note that this is a function of the population distributions. We will want to produce estimates  $\hat{r}(x)$ .

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- Note that these are just **conditional expectations**. Define  $Y$  to be the loan amount,  $X = 1$  to indicate a man, and  $X = 0$  to indicate a woman and then we have:

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- Notice here that since  $X$  can only take on two values, 0 and 1, then these two conditional means completely summarize the CEF.

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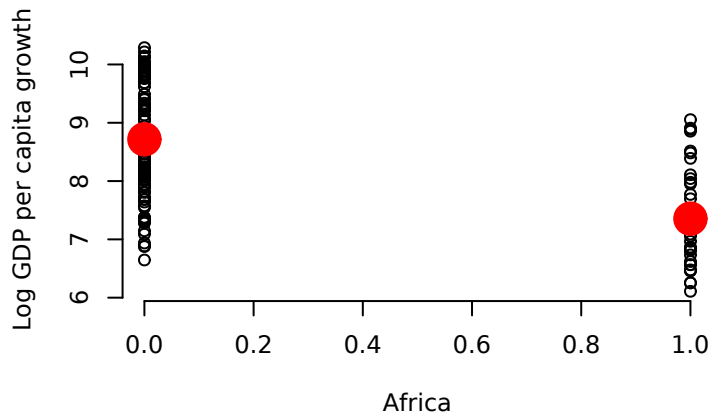
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- This is very straightforward: estimate the mean of  $Y$  conditional on  $X$  by just estimating the means within each group of  $X$ .



## Binary covariate example CEF plot



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- We want to extend the regression idea to the case of multiple  $X$  variables, but we will start this week with the simple bivariate case where we have a single  $X$

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## Fun With Salmon

Bennett, Baird, Miller and Wolford. (2009). "Neural correlates of interspecies perspective taking in the post-mortem Atlantic Salmon: an argument for multiple comparisons correction."



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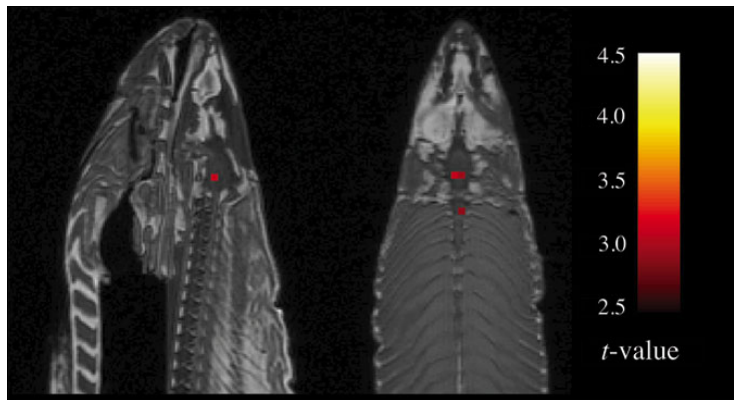
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## Results



“Several active voxels were discovered in a cluster located within the salmon’s brain cavity. The size of this cluster was  $81 \text{ mm}^3$  with a cluster-level significance of  $p = .001$ .”



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Questions?

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- Let's take a look at some data on education and income from the American National Election Study
- We use two variables:
  - ▶  $Y$ : income
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- Goal is to characterize the conditional expectation  $E[Y|X]$ , i.e. how average income varies with education level

# Nonparametric Regression with Discrete $X$



# Nonparametric Regression with Discrete $X$

educ: Respondent's education:

- 1. 8 grades or less and no diploma or
- 2. 9-11 grades
- 3. High school diploma or equivalency test
- 4. More than 12 years of schooling, no higher degree
- 5. Junior or community college level degree (AA degrees)
- 6. BA level degrees; 17+ years, no postgraduate degree
- 7. Advanced degree

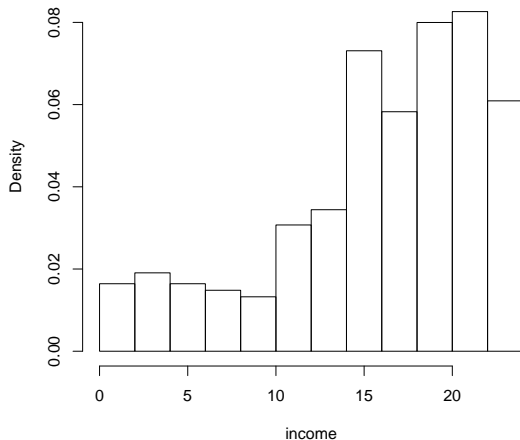
# Nonparametric Regression with Discrete $X$

income: Respondent's family income:

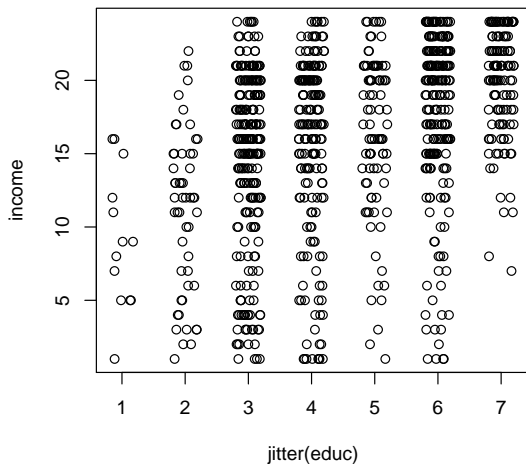
- 1. None or less than \$2,999
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- ⋮
- 17. \$35,000-\$39,999
- 18. \$40,000-\$44,999
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- 23. \$90,000-\$104,999
- 24. \$105,000 and over

# Marginal Distribution of $Y$

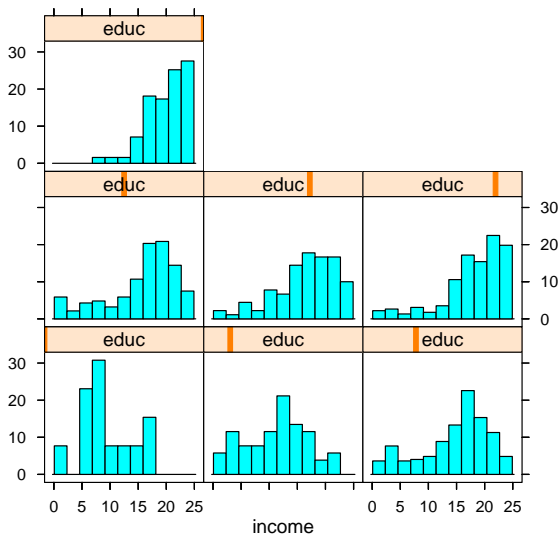
Histogram of income



# Income and Education



# Distribution of income given education $p(y|x)$



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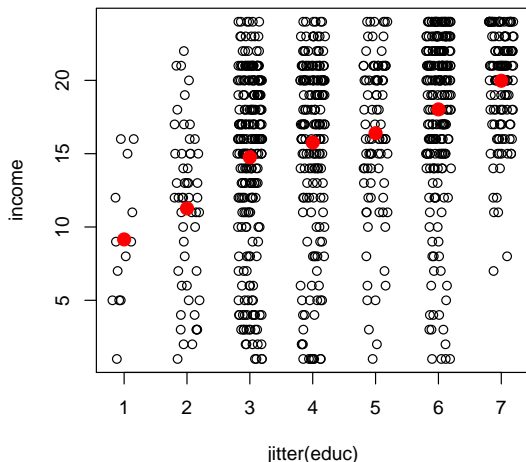
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- Here our  $X$  variable education has a small number of levels (7) and there are a reasonable number of observations in each level
- In situations like this we can estimate  $E[Y|X = x]$  as the sample mean of  $Y$  at each level of  $x \in X$  (just like the binary case)

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- But what do we do when  $X$  is continuous and has many values?

# Nonparametric Regression with Continuous $X$

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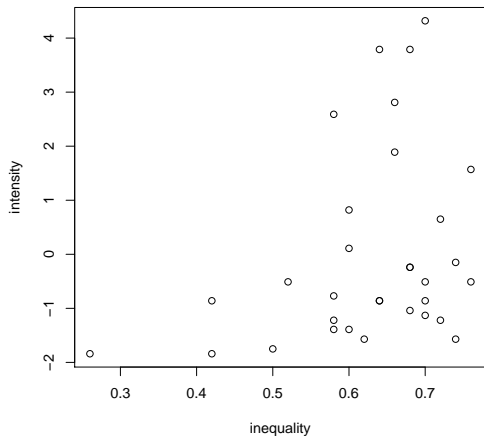
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- Around 11,000 peasants were killed by Romanian military

# Nonparametric Regression with Continuous $X$



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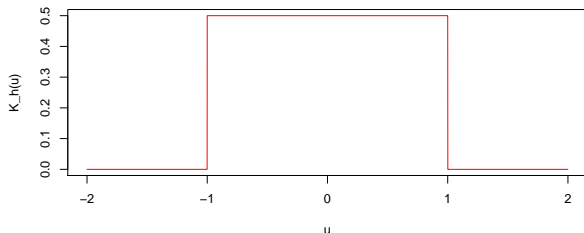
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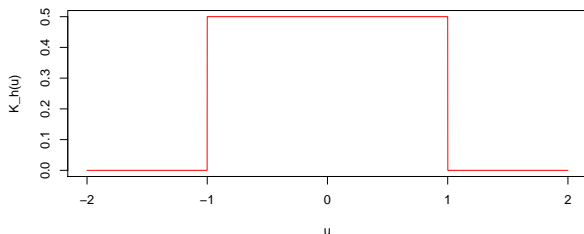
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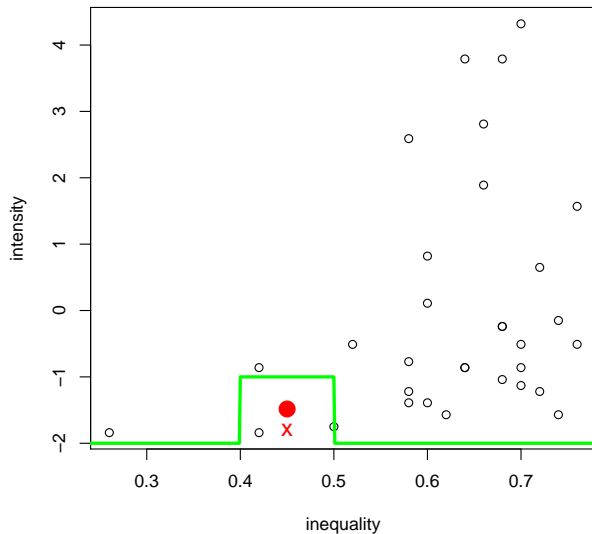
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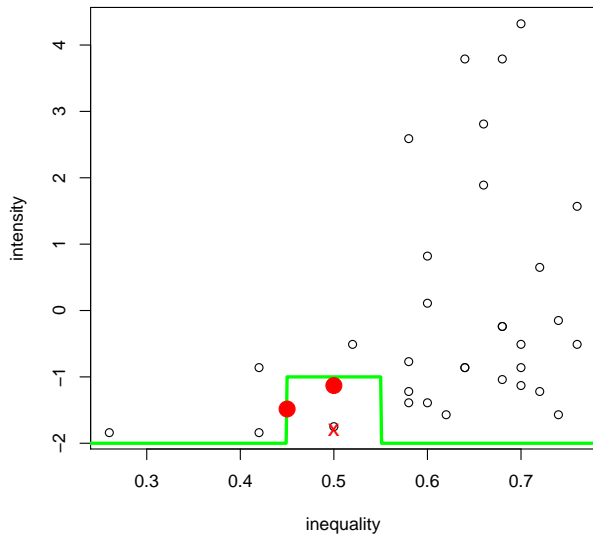
- This gives the **uniform kernel regression**:

$$\hat{E}[Y|X = x_0] = \frac{\sum_{i=1}^N K_h((X_i - x_0)/h) Y_i}{\sum_{i=1}^N K_h((X_i - x_0)/h)} \text{ where } K_h(u) = \frac{1}{2} \mathbf{1}_{\{|u| \leq 1\}}$$

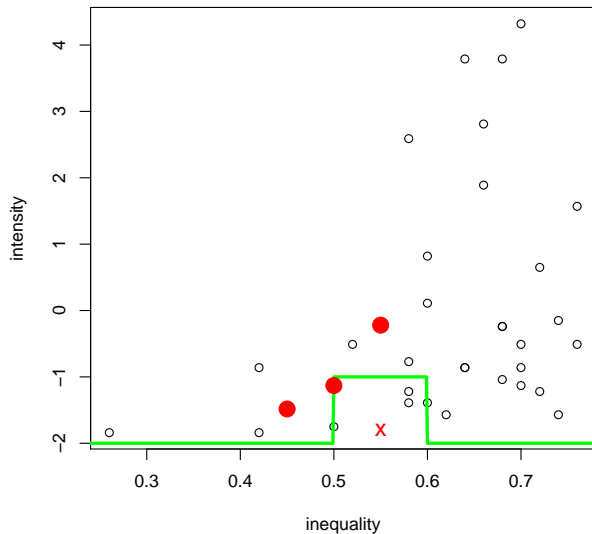
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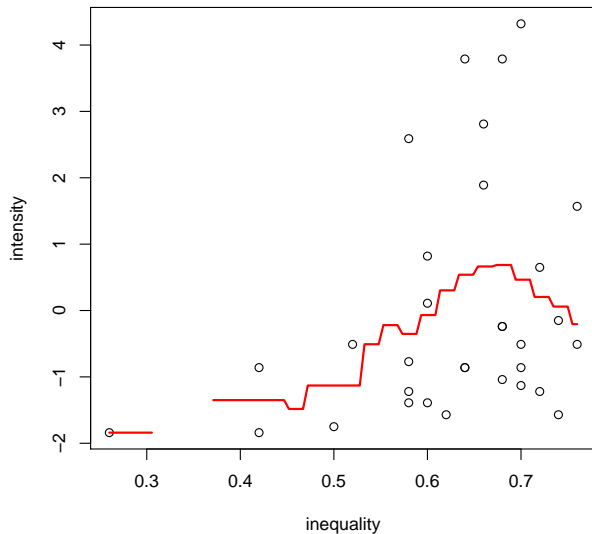
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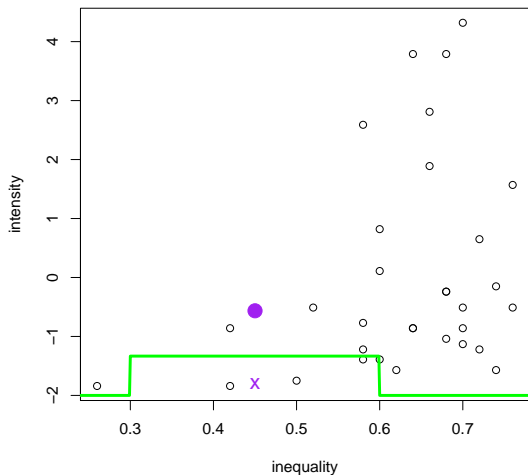
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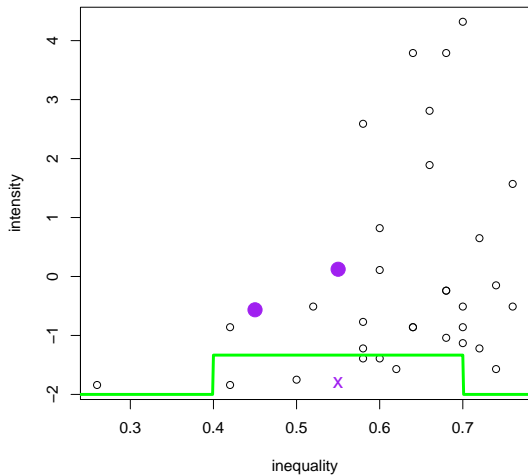
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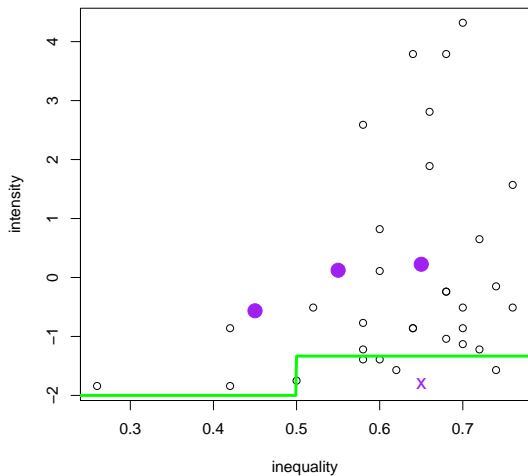
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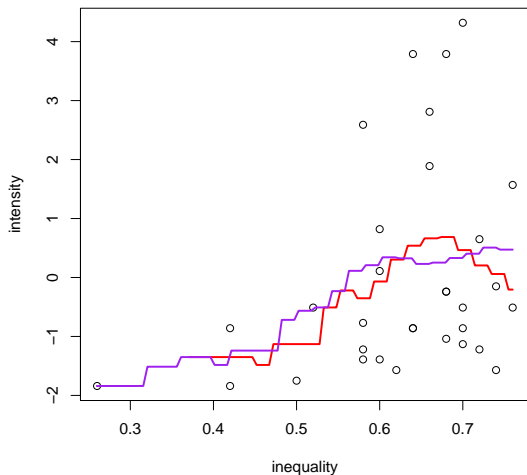


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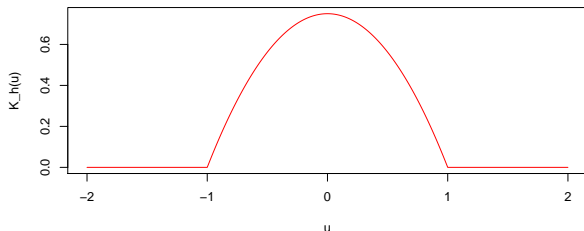
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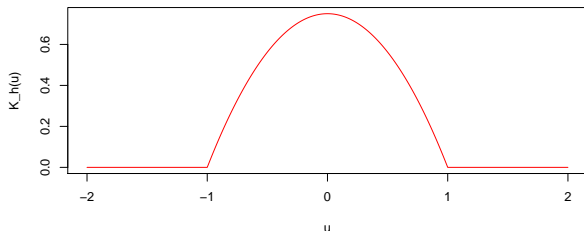
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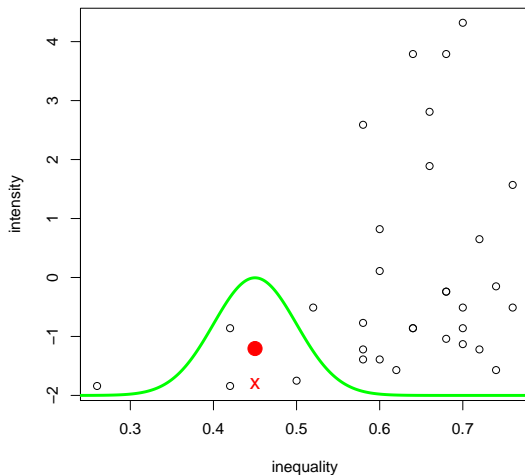
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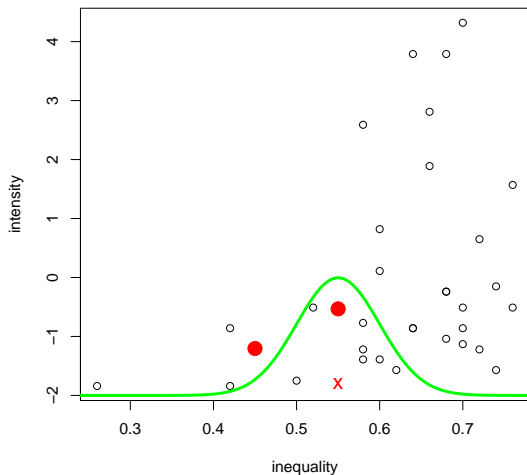
- ② compute weighted average of the observed  $y$  points that have  $x$  values in the bandwidth interval  $[x_0 - h, x_0 + h]$  e.g.

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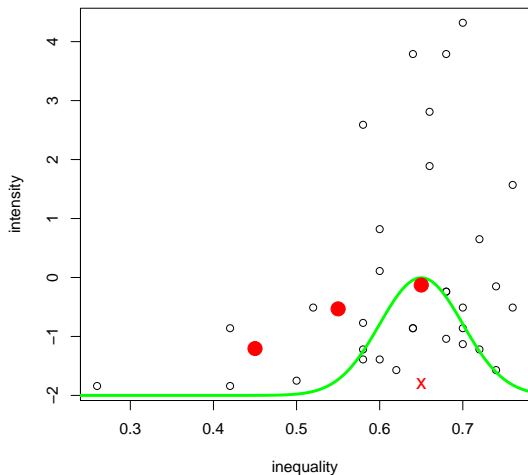


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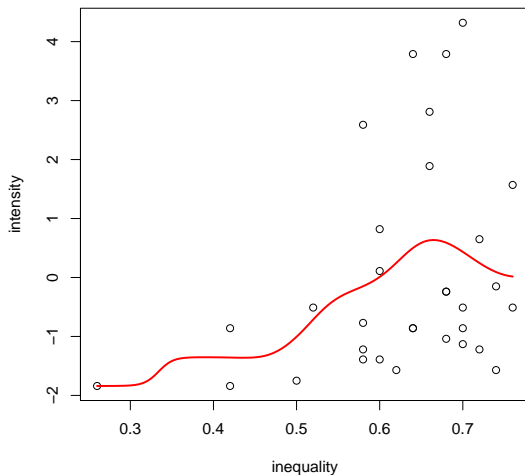




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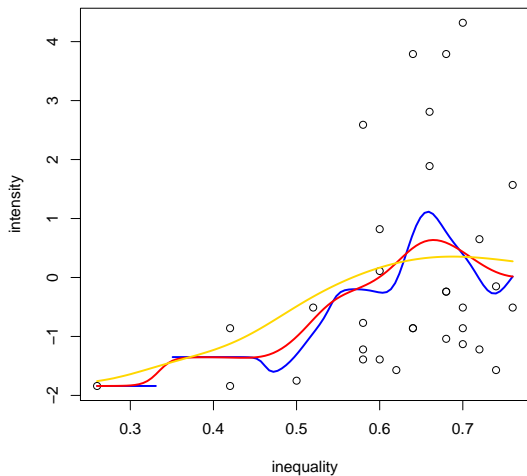


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  - ▶ A very **flexible estimator** allows the shape of the function to vary (e.g. a kernel regression with a small bandwidth)
  - ▶ A very **inflexible estimator** restricts the shape of the function to a particular form (e.g. a kernel regression with a very wide bandwidth)

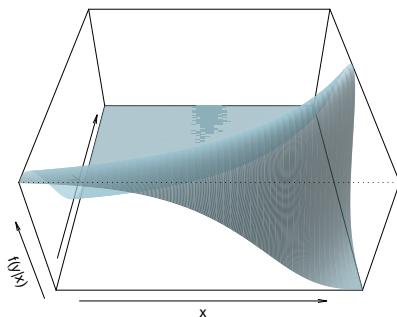
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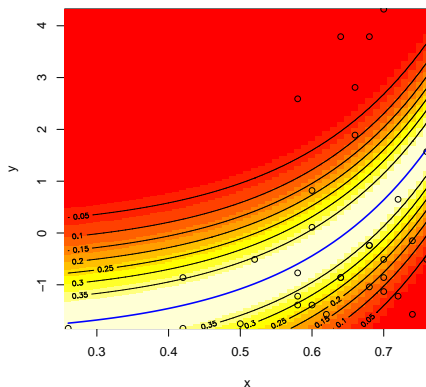
- Let's conduct a simulation experiment to actually see the tradeoff
- Suppose we have the following population distribution:



# Hypothetical True Distribution

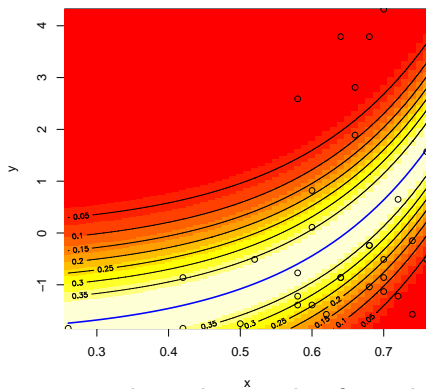
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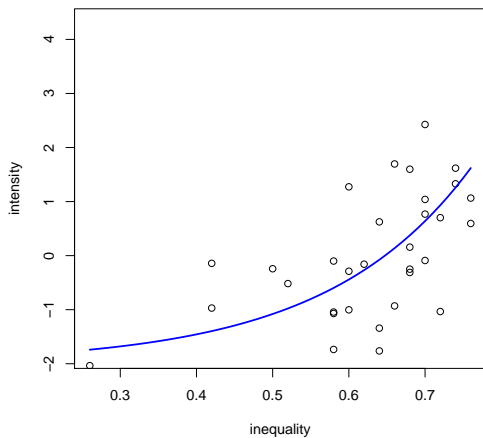
# Hypothetical True Distribution

- Another way of representing the same population distribution:



- From this distribution we draw thousands of simulated data sets.

# An Example of Simulated Data Set





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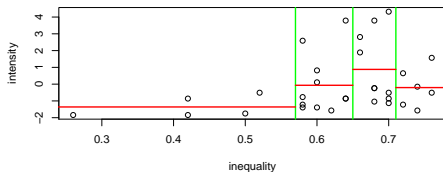
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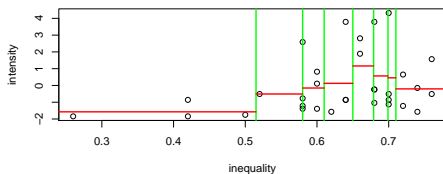
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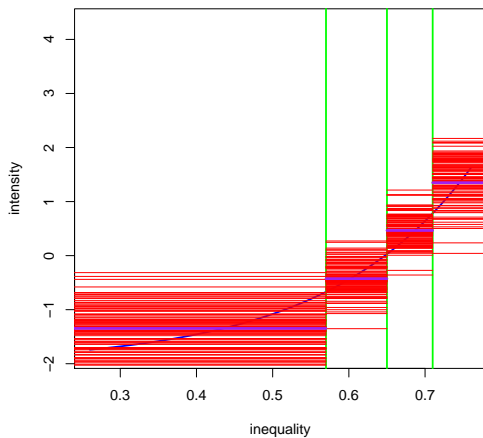
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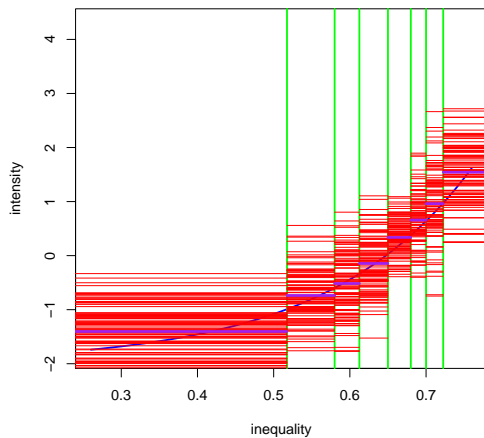
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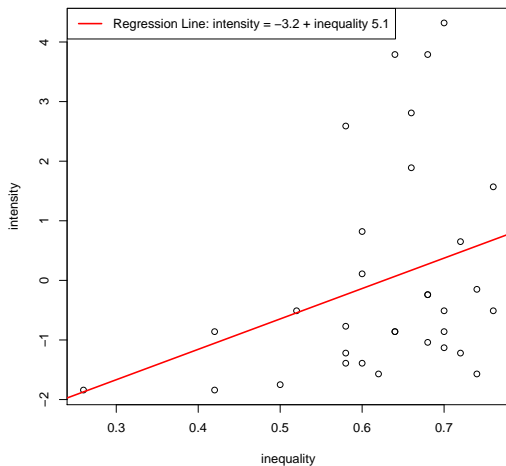
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Figure: 'If I fits, I sits'

Linear regression always returns a **line** regardless of the data.

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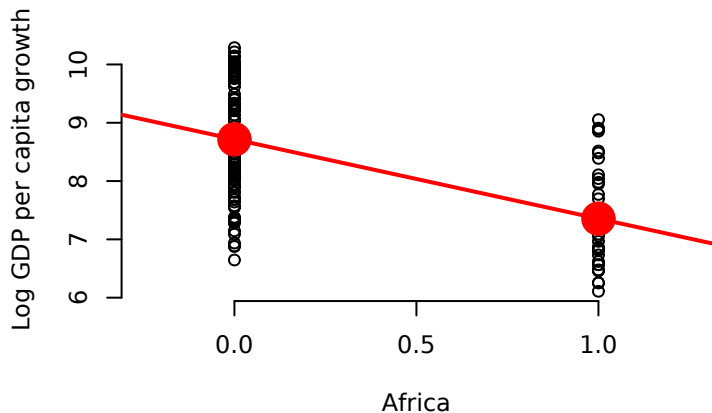
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$$\beta_1 = E[Y|X = 1] - E[Y|X = 0]$$
- Thus, we can read off the difference in means between two groups as the slope coefficient on a linear regression

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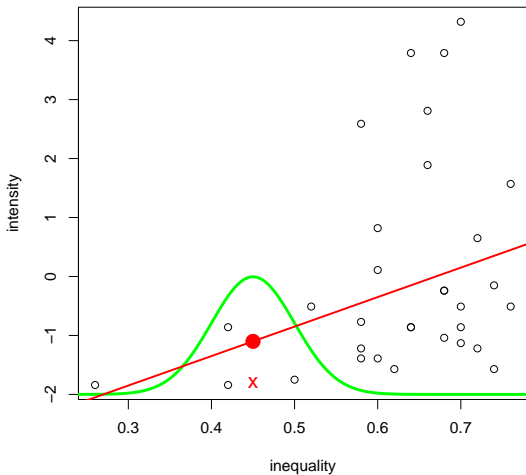
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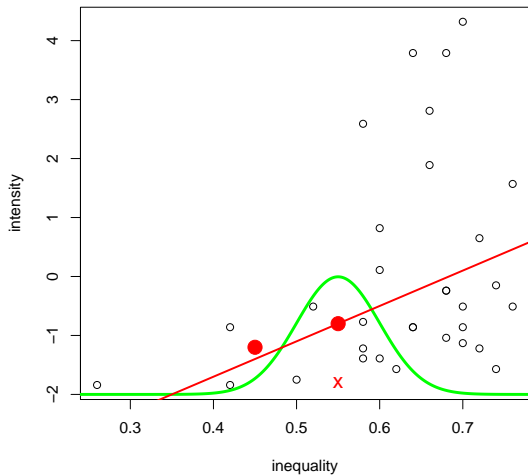
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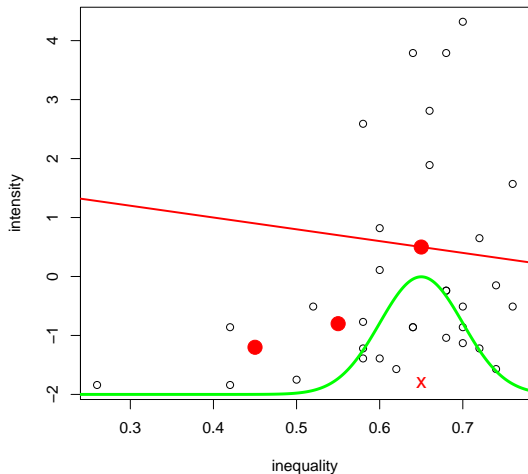
# Weighted Local Linear Regressions



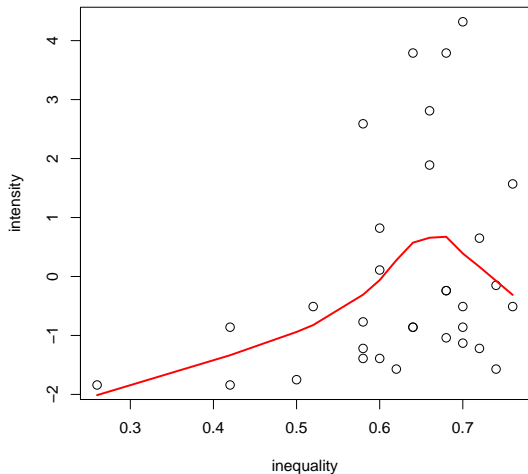
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- Need to estimate them in our samples! But how?

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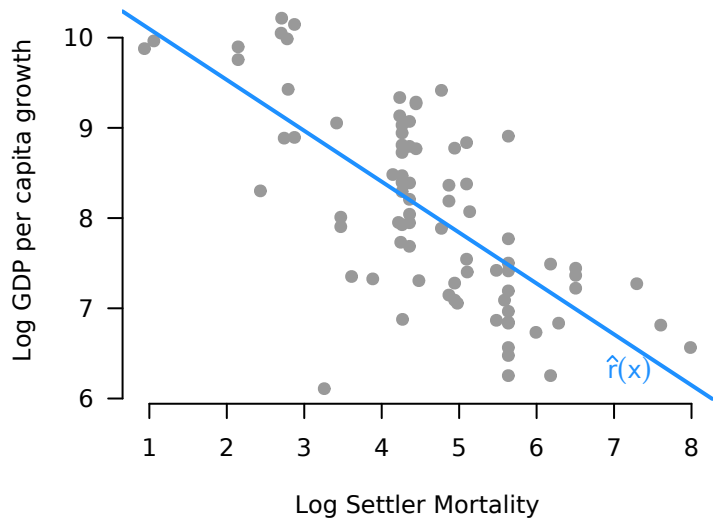
$$Y_i = \beta_0 + \beta_1 X_i + u_i$$

- Now, suppose we have some estimates of the slope,  $\hat{\beta}_1$ , and the intercept,  $\hat{\beta}_0$ . Then the fitted or sample regression line is

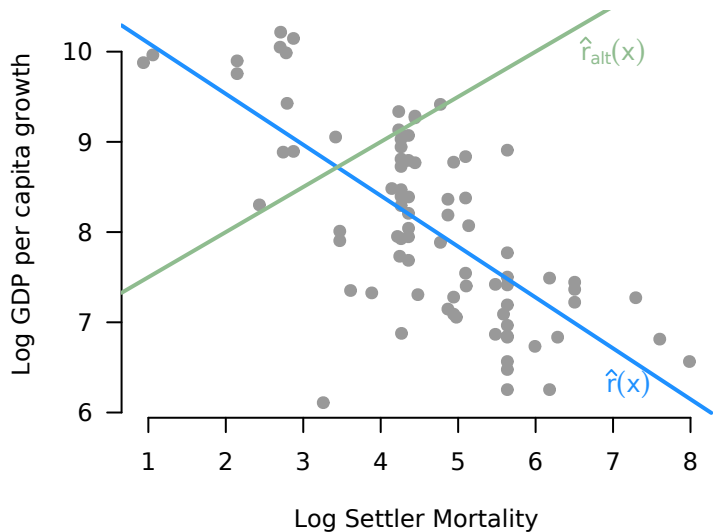
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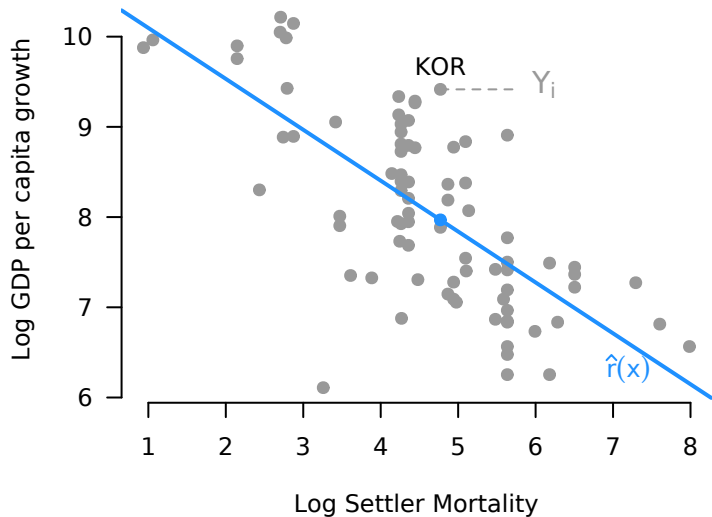
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## Definition (Residual)

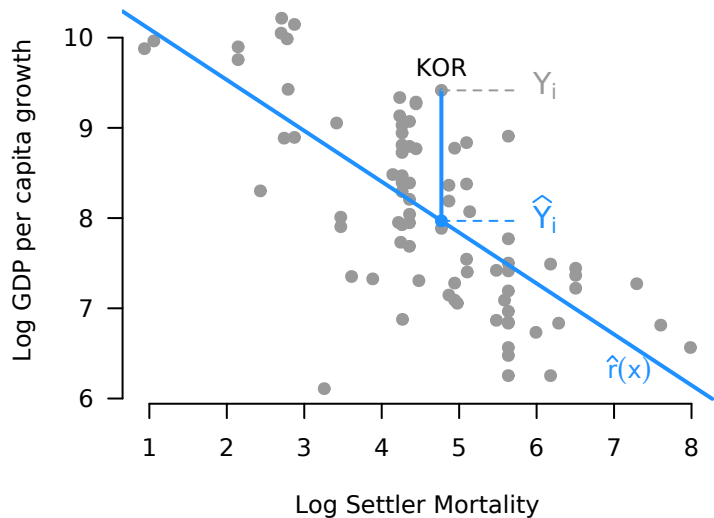
The **residual** is the difference between the actual value of  $Y_i$  and the predicted value,  $\hat{Y}_i$ :

$$\hat{u}_i = Y_i - \hat{Y}_i = Y_i - \hat{\beta}_0 - \hat{\beta}_1 X_i$$

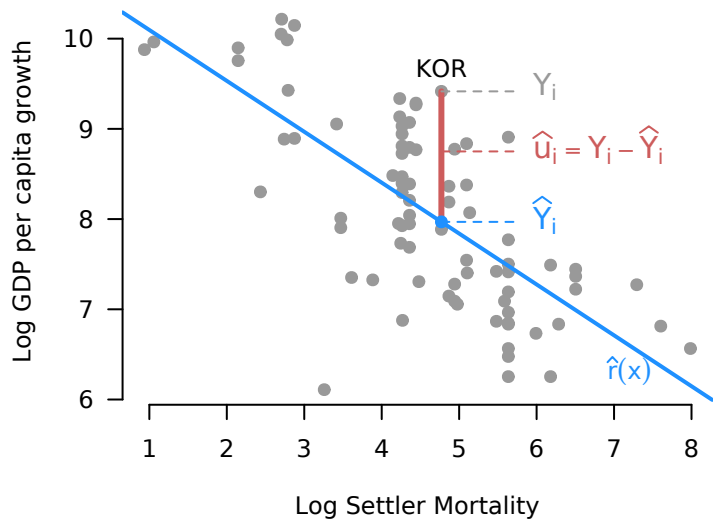
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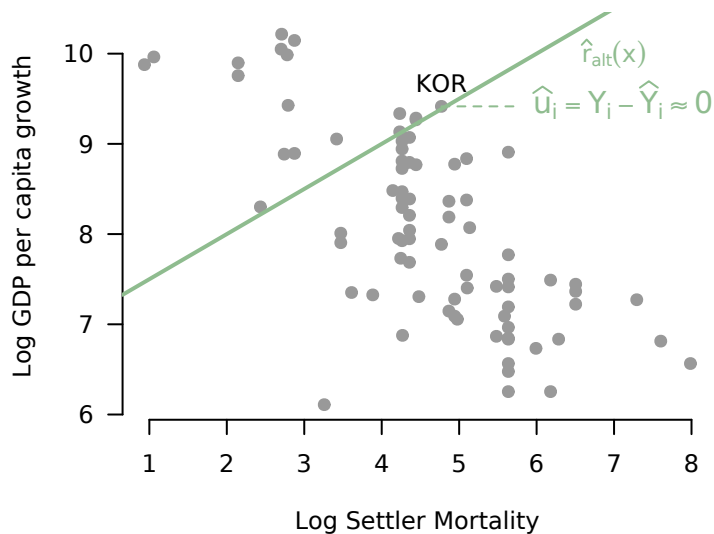


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## Why not this line?



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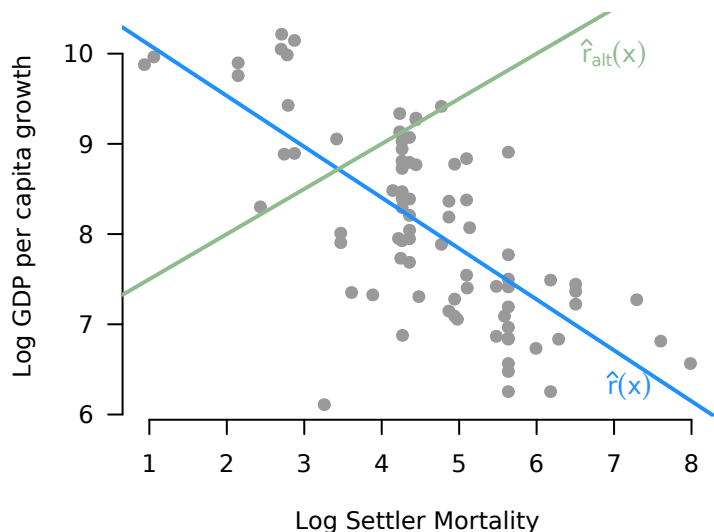
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# Which is better at minimizing residuals?





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- Least squares is optimal in a certain sense that we'll see in the coming weeks

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  - ▶ We can make the model more flexible, even in a linear framework (e.g. we can add polynomials, use log transformations, etc.)

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  - Forming rejection regions
  - P-values
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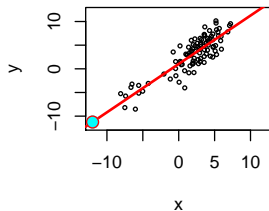
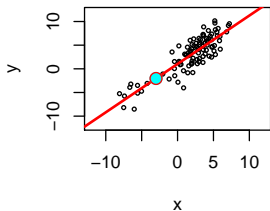
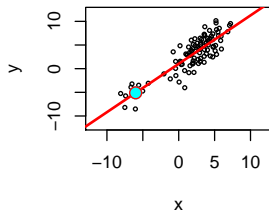
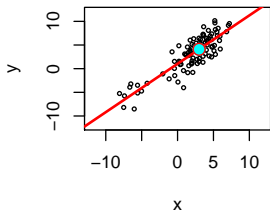
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- While the line is defined over all regions of the data we may be concerned about:
  - ▶ interpolation
  - ▶ extrapolation
  - ▶ predicting in ranges of  $X$  with sparse data

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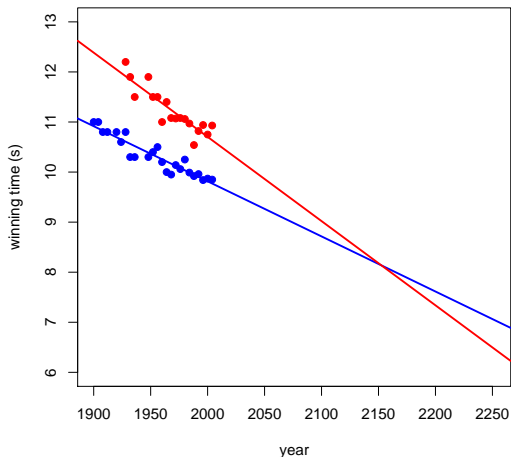
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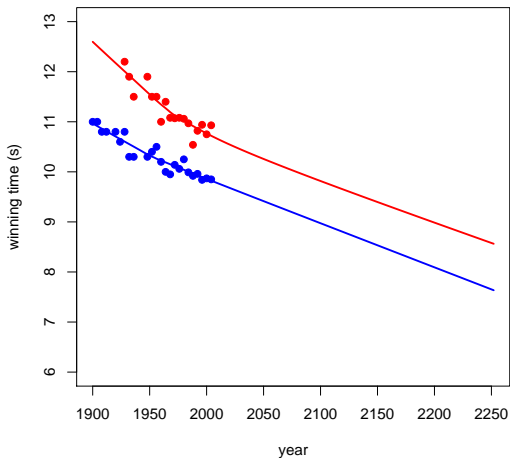
# Tatem et al. Extrapolation



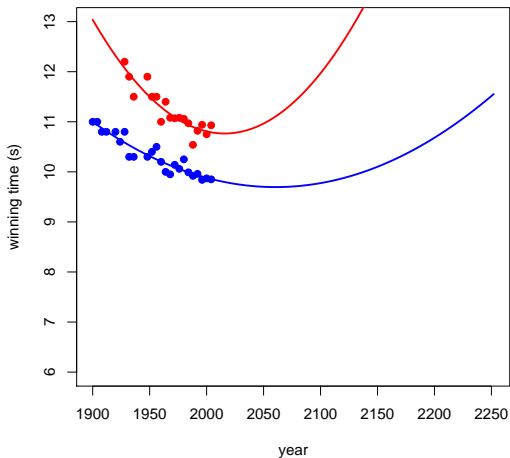
Tatem et al.'s predictions. Men's times are in blue, women's times are in red.



# Alternate Models Fit Well, Yield Different Predictions



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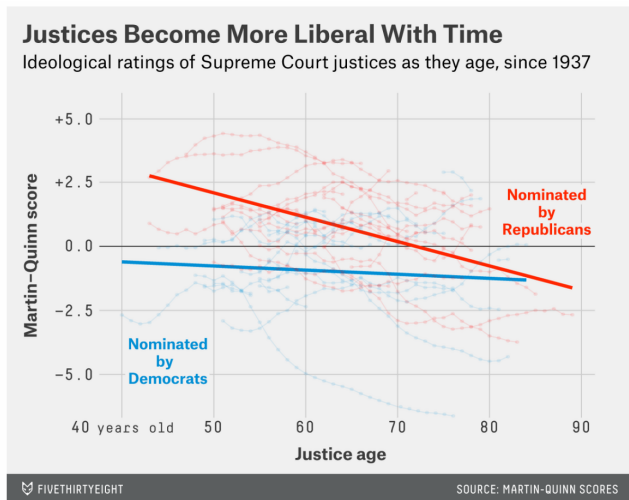
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- Next semester we will talk about how this problem gets much harder in high dimensions



# A More Subtle Example

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the signal  
and the noise  
why so many  
predictions fail—  
but some don't

**Nate Silver**  @NateSilver538 · Oct 5

So, basically, John Roberts is going to be Ruth Bader Ginsburg by 2036.

[53eig.ht/1Gsl2u6](https://53eig.ht/1Gsl2u6)



## Supreme Court Justices Get More Liberal As They Get Older

The Supreme Court justices are back from vacation. They've picked up their robes from the cleaners — Alito's had a pesky mustard stain — and are ...

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- Always think about where we have data and what we are using to build our claims
- Summary: 'prediction is hard, especially about the future'

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- For now, it is safest to treat  $\beta$  as a purely descriptive/predictive quantity

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- Reading:
  - ▶ Fox Chapter 6.1
  - ▶ Optional: Imai 4.2

## **Iterated learning: Intergenerational knowledge transmission reveals inductive biases**

**MICHAEL L. KALISH**

*University of Louisiana, Lafayette, Louisiana*

**THOMAS L. GRIFFITHS**

*University of California, Berkeley, California*

AND

**STEPHAN LEWANDOWSKY**

*University of Western Australia, Perth, Australia*

Cultural transmission of information plays a central role in shaping human knowledge. Some of the most complex knowledge that people acquire, such as languages or cultural norms, can only be learned from other people, who themselves learned from previous generations. The prevalence of this process of “iterated learning” as a mode of cultural transmission raises the question of how it affects the information being transmitted. Analyses of iterated learning utilizing the assumption that the learners are Bayesian agents predict that this process should converge to an equilibrium that reflects the inductive biases of the learners. An experiment in iterated function learning with human participants confirmed this prediction, providing insight into the consequences of intergenerational knowledge transmission and a method for discovering the inductive biases that guide human inferences.

Images on following slides courtesy of Tom Griffiths



# Fun with Linearity



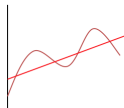
# The Design

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**data**



**hypotheses**

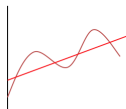


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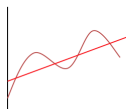
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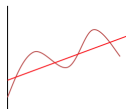
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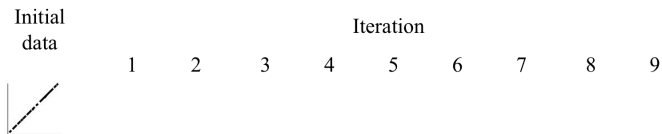


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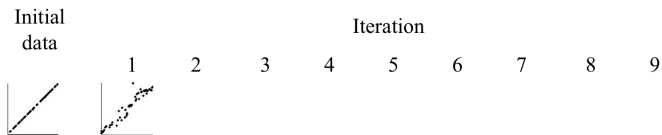


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- Predictions are data for the next learner

# Results

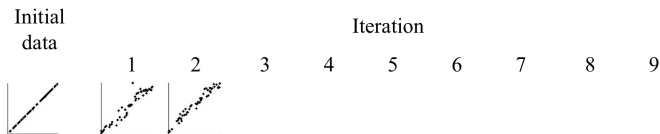


# Results

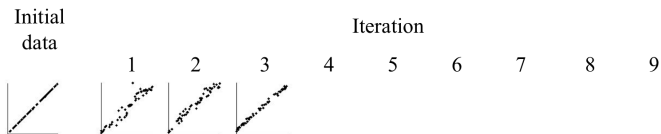




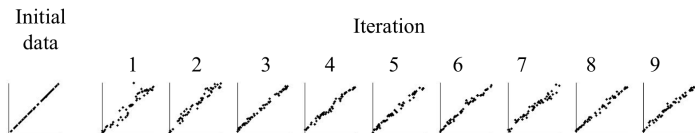
# Results



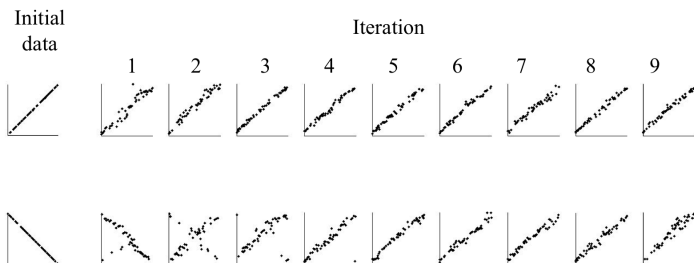
# Results



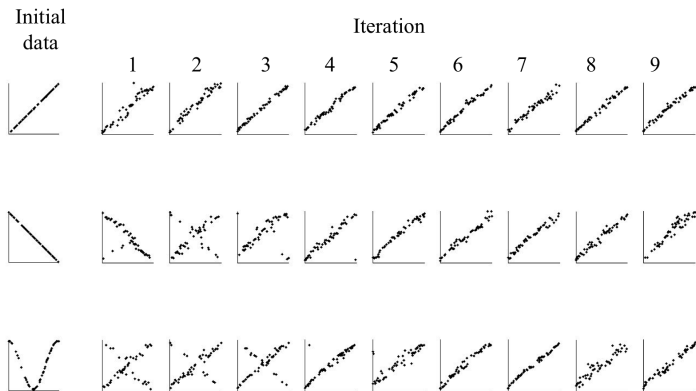
# Results



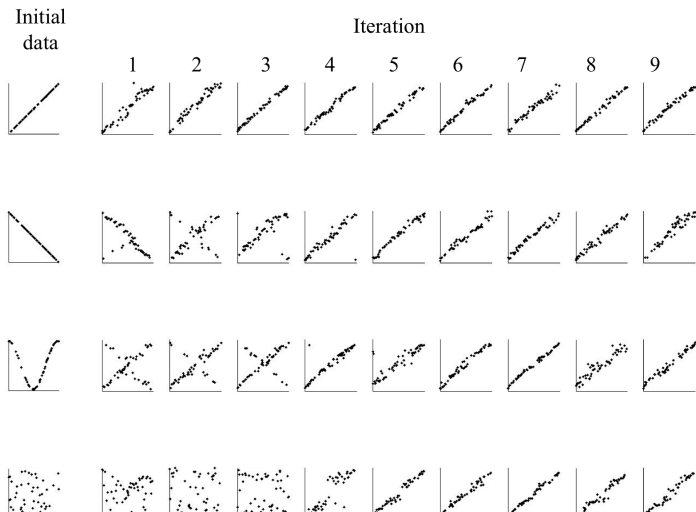
# Results



# Results



# Results



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