

# SOC-5811 Week 4: Linear regression

Nick Graetz

University of Minnesota, Department of Sociology

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1/73



# LOAD DATA

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- ▶ We apply functions to objects: **function(object)**.
- ▶ All objects have a **class**.
- ▶ All functions take inputs of certain classes and return outputs of certain classes.
  
- ▶ **Scripts are run in computing environments.**



# SET UP FILEPATHS

- ▶ Assign my filepath to an object called “dropbox”
- ▶ Test different basic R functions

```
dropbox <- 'C:/Users/ngraetz/Dropbox/'  
class(dropbox)
```

```
## [1] "character"
```

```
length(dropbox)
```

```
## [1] 1
```

```
nchar(dropbox)
```

```
## [1] 25
```



# SET UP FILEPATHS

```
sum(dropbox)
```

```
## Error in sum(dropbox): invalid 'type' (character) of
```



# SET UP FILEPATHS

- I'm going to set up a few filepaths.

```
fig_dir <- paste0(dropbox,  
                  'Minnesota/repos/soc5811/figures/')  
data_dir <- paste0(dropbox,  
                   'Minnesota/repos/soc5811/data/')
```

# LOAD DATA

- ▶ We are going to look at population and housing data from the 2000/2010 Census.

```
census <- read_dta(paste0(data_dir,  
                           'state_pophouse.dta'))
```

# LOAD DATA

```
class(census)
```

```
## [1] "tbl_df"      "tbl"        "data.frame"
```

```
dim(census)
```

```
## [1] 51  9
```

```
names(census)
```

```
## [1] "state"      "statefp"    "a00aa2000" "a00aa2010" "a41aa2000" "a41aa2010"  
## [7] "pctpop"     "pcthouse"   "onepct"
```

```
head(census)
```

```
## # A tibble: 6 x 9  
##   state statefp a00aa2000 a00aa2010 a41aa2000 a41aa2010 pctpop pcthouse onepct  
##   <chr>   <chr>         <dbl>      <dbl>      <dbl>      <dbl>    <dbl>    <dbl>    <dbl>  
## 1 Alabama 01           4447100    4779736    1963711    2171853     7.48     10.6      0  
## 2 Alaska  02           626932     710231     260978     306967     13.3     17.6      1  
## 3 Arizona 04           5130632    6392017    2189189    2844526     24.6     29.9      2  
## 4 Arkans~ 05           2673400    2915918    1173043    1316299     9.07     12.2      3  
## 5 Califo~ 06           33871648   37253956   12214549   13680081     9.99     12.0      4  
## 6 Colora~ 08           4301261    5029196    1808037    2212898    16.9     22.4      5
```

7/73



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# EXAMINE DATA

We can use different functions like `select()` and `slice()` to look at specific rows and columns:

```
census %>%  
  select(state, a00aa2000) %>%  
  slice(1:5)
```

```
## # A tibble: 5 x 2  
##   state      a00aa2000  
##   <chr>      <dbl>  
## 1 Alabama    4447100  
## 2 Alaska     626932  
## 3 Arizona    5130632  
## 4 Arkansas   2673400  
## 5 California 33871648
```

# EXAMINE DATA

We can use other packages like “data.table” with different functions:

```
census <- as.data.table(census)
class(census)
```

```
## [1] "data.table" "data.frame"
```

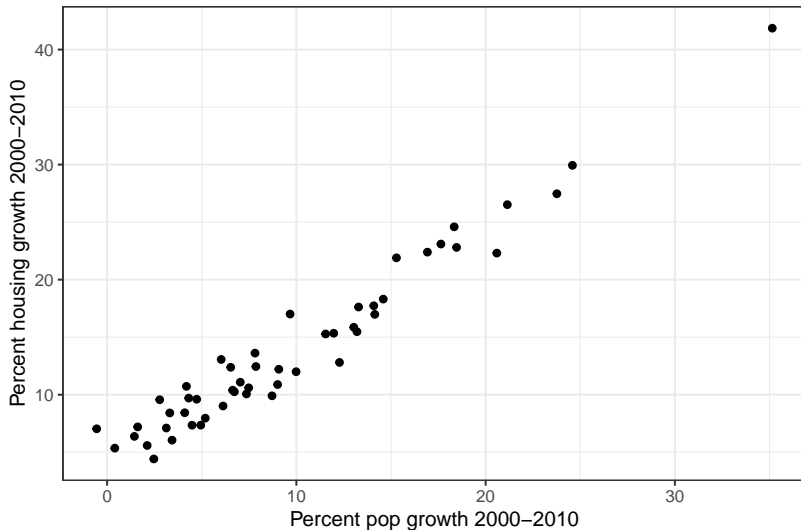
```
census[1:5, c('state', 'a00aa2000')]
```

```
##           state a00aa2000
##      <char>    <num>
## 1:  Alabama    4447100
## 2:   Alaska    626932
## 3:  Arizona    5130632
## 4: Arkansas    2673400
## 5: California  33871648
```

# EXAMINE DATA

```
plot <- ggplot(data=census,  
               aes(x=pctpop,  
                   y=pcthouse)) +  
  geom_point() +  
  labs(x='Percent pop growth 2000-2010',  
       y='Percent housing growth 2000-2010') +  
  theme_bw()
```

# EXAMINE DATA

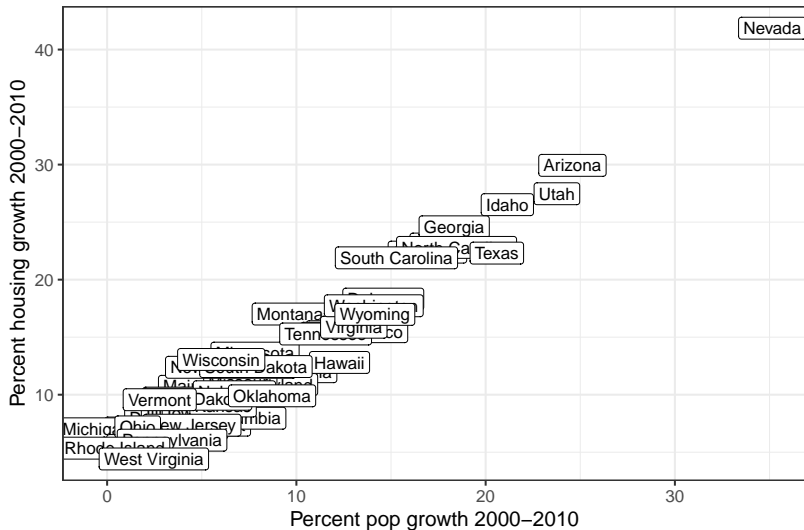




# EXAMINE DATA

```
plot <- ggplot(data=census,  
               aes(x=pctpop,  
                   y=pcthouse,  
                   label=state)) +  
  geom_label(size=3) +  
  labs(x='Percent pop growth 2000-2010',  
       y='Percent housing growth 2000-2010') +  
  theme_bw()
```

# EXAMINE DATA



# POPULATION REGRESSION FUNCTIONS

Let's think about creating a **model** for housing growth:

$$pcthouse = f(pctpop)$$

- What is a model?

# POPULATION REGRESSION FUNCTIONS

Let's think about creating a **model** for housing growth:

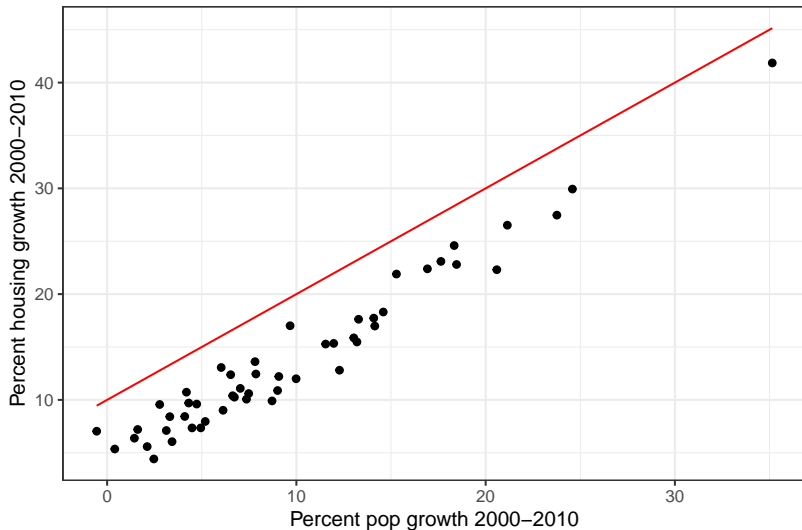
$$pcthouse = 10 + pctpop$$

- Models are defined by *coefficients* (or more generally, *parameters*).

# EXAMINE DATA

```
census <- census %>%  
  mutate(pcthouse_mod1 = 10 + pctpop)
```

# EXAMINE DATA



# POPULATION REGRESSION FUNCTIONS

Let's think about creating a **model** for housing growth:

$$pcthouse = 1.5 \times pctpop$$

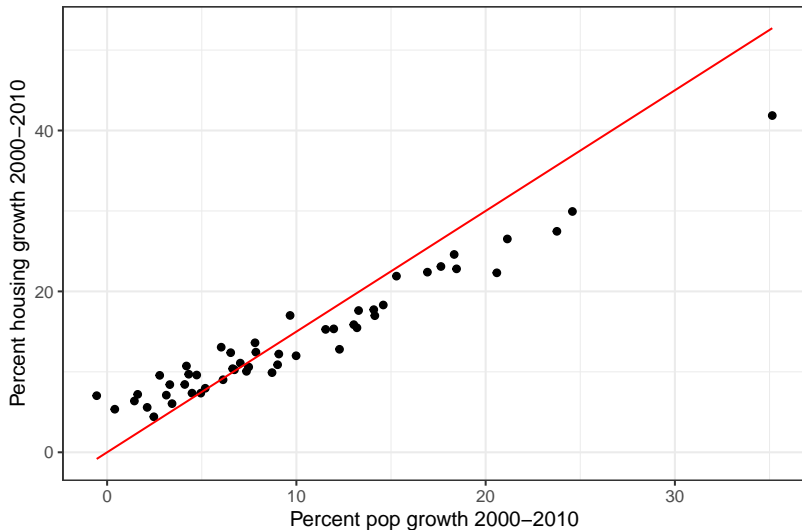


# EXAMINE DATA

```
census <- census %>%  
  mutate(pcthouse_mod1 = 1.5 * pctpop)
```



# EXAMINE DATA



20/73



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# POPULATION REGRESSION FUNCTIONS

- ▶ How do I pick a good model?
- ▶ What makes a model good?
- ▶ What is my goal?

$$pcthouse = f(pctpop)$$

# POPULATION REGRESSION FUNCTIONS

Let's think about creating a model for housing growth:

$$pcthouse = f(pctpop)$$

$$pcthouse_i = \beta_0 + \beta_1 pctpop_i + \epsilon_i$$

# POPULATION REGRESSION FUNCTIONS

## Fitting a linear regression with data:

```
model <- lm(pcthouse~pctpop,  
            data=census)  
summary(model)
```

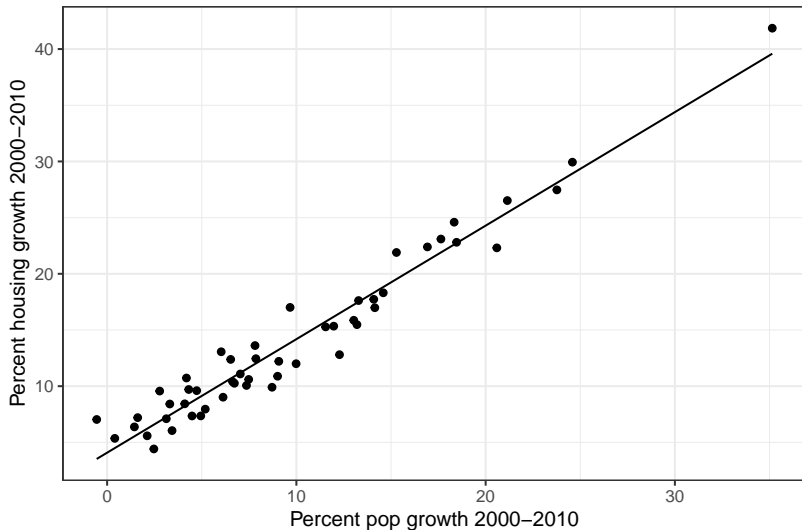
```
##  
## Call:  
## lm(formula = pcthouse ~ pctpop, data = census)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max   
## -3.6830 -1.3132 -0.1364  1.2039  3.5126   
##  
## Coefficients:  
##              Estimate Std. Error t value Pr(>|t|)      
## (Intercept)  4.08125    0.40793   10.01 1.98e-13 ***  
## pctpop       1.01030    0.03371   29.97 < 2e-16 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 1.719 on 49 degrees of freedom  
## Multiple R-squared:  0.9483, Adjusted R-squared:  0.9472   
## F-statistic: 898.1 on 1 and 49 DF,  p-value: < 2.2e-16
```

# MAKING PREDICTIONS

```
census <- census %>%  
  mutate(pcthouse_pred=predict(model))  
census %>%  
  select(state,pctpop,pcthouse,pcthouse_pred) %>%  
  head()
```

	state	pctpop	pcthouse	pcthouse_pred
	<char>	<num>	<num>	<num>
## 1:	Alabama	7.479841	10.59942	11.63810
## 2:	Alaska	13.286768	17.62179	17.50481
## 3:	Arizona	24.585373	29.93515	28.91974
## 4:	Arkansas	9.071520	12.21234	13.24616
## 5:	California	9.985662	11.99825	14.16972
## 6:	Colorado	16.923758	22.39230	21.17924

# MAKING PREDICTIONS



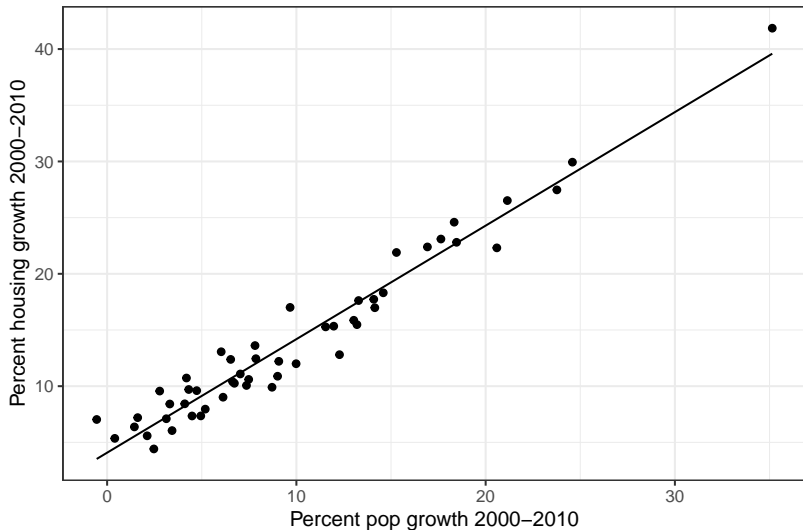
# MODEL COEFFICIENTS

- ▶ What does it mean to “fit” a regression model?
- ▶ How did R come up with the coefficients 4.08 and 1.01?

```
summary(model)
```

```
##
## Call:
## lm(formula = pcthouse ~ pctpop, data = census)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.6830 -1.3132 -0.1364  1.2039  3.5126
##
## Coefficients:
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## (Intercept)   4.08125    0.40793   10.01 1.98e-13 ***
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## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
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## Residual standard error: 1.719 on 49 degrees of freedom
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## F-statistic: 898.1 on 1 and 49 DF,  p-value: < 2.2e-16
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# MAKING PREDICTIONS



27/73



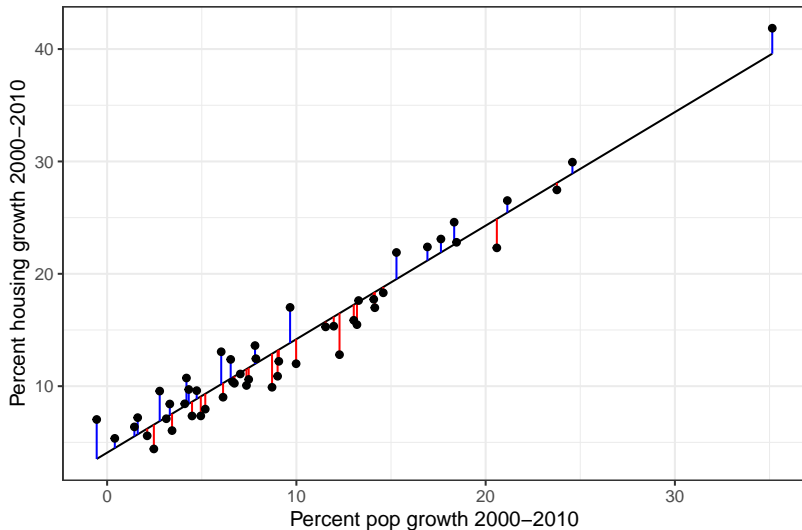
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# MAKING PREDICTIONS



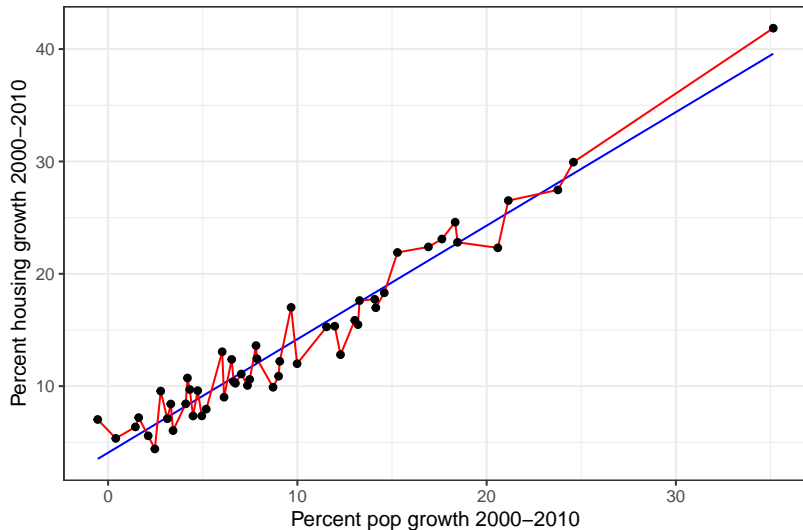
# USING MODELS TO PREDICT

Just looking *within* my sample... why is my model always wrong?

```
census %>%  
  select(state, pctpop, pcthouse, pcthouse_pred) %>%  
  head()
```

##	state	pctpop	pcthouse	pcthouse_pred
##	<char>	<num>	<num>	<num>
## 1:	Alabama	7.479841	10.59942	11.63810
## 2:	Alaska	13.286768	17.62179	17.50481
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# USING MODELS TO PREDICT



30/73



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# COEFFICIENTS

$$pcthouse_i = \beta_0 + \beta_1 pctpop_i + \epsilon_i$$

$$pcthouse_i = 4.08 + 1.01 pctpop_i + \epsilon_i$$

- Coefficients represent **average comparisons**.

# COEFFICIENTS

$$pcthouse_i = \beta_0 + \beta_1 pctpopt_i + \epsilon_i$$

$$pcthouse_i = 4.08 + 1.01 pctpopt_i + \epsilon_i$$

- ▶ Coefficients represent **average comparisons**.
- ▶ Interpreting the coefficient on *pctpopt* (e.g.,  $x$ ):

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- ▶ Coefficients represent **average comparisons**.
- ▶ Interpreting the coefficient on *pctpop* (e.g.,  $x$ ):
  - ▶ On average, a 1-point increase in  $x$  is associated with a 1.01-point increase in  $y$ .

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- ▶ Interpreting the coefficient on *pctpop* (e.g.,  $x$ ):
  - ▶ On average, a 1-point increase in  $x$  is associated with a 1.01-point increase in  $y$ .
  - ▶ Across all values of  $x$ , the average difference in  $y$  at  $x$  and  $x+1$  is 1.01.

# COEFFICIENTS

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- ▶ Coefficients represent **average comparisons**.
- ▶ Interpreting the coefficient on *pctpopt* (e.g.,  $x$ ):
  - ▶ On average, a 1-point increase in  $x$  is associated with a 1.01-point increase in  $y$ .
  - ▶ Across all values of  $x$ , the average difference in  $y$  at  $x$  and  $x+1$  is 1.01.
  - ▶ The slope of the predicted line of  $y$  across all values of  $x$  is 1.01.



# USING MODELS TO COMPARE

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- ▶ Regression coefficients can *sometimes* be interpreted as effects.



# USING MODELS TO COMPARE

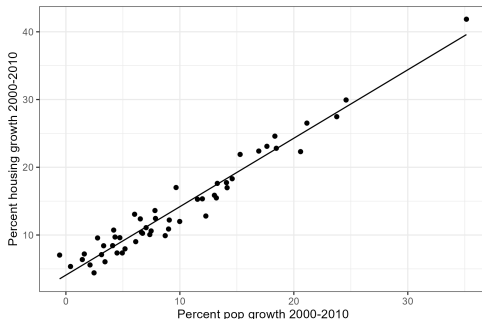
- ▶ Regression is a mathematical tool for making predictions.
- ▶ Regression coefficients can *sometimes* be interpreted as effects.
- ▶ Regression coefficients can *always* be interpreted as average comparisons.



# BUILDING MODELS

What can we do with this model?

1. Generalizing from sample to population.
2. Measurement.
3. Forecasting.
4. Causal inference.

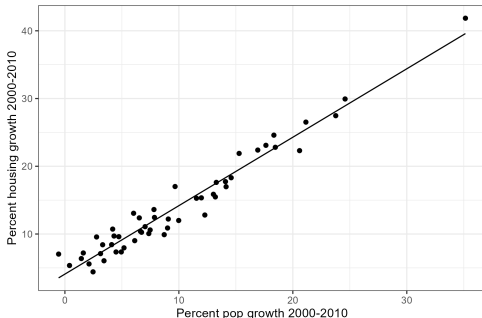


# BUILDING MODELS

What can we do with this model?

1. **Generalizing from sample to population:**

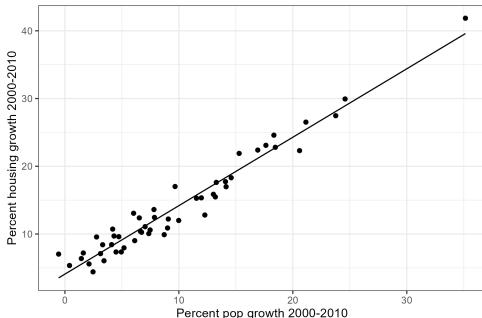
Is this coefficient the same one I would estimate with the entire population?



# BUILDING MODELS

What can we do with this model?

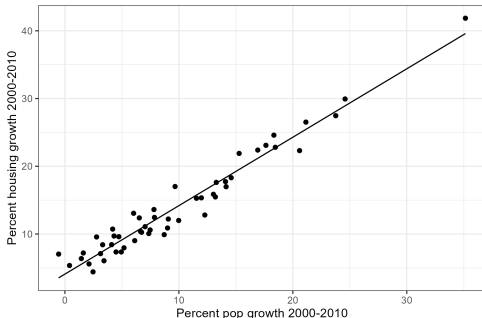
1. Generalizing from sample to population
2. **Measurement:** Can I generalize to all types of housing growth?



# BUILDING MODELS

What can we do with this model?

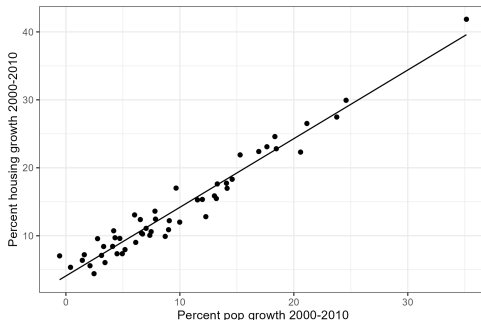
1. Generalizing from sample to population
2. Measurement
3. **Forecasting:** Can I use this model to predict out-of-sample?



# BUILDING MODELS

What can we do with this model?

1. Generalizing from sample to population
2. Measurement
3. Forecasting
4. **Causal inference:** Can I say pop growth **causes** housing growth?





# REVIEW

- ▶ How do we calculate linear regression coefficients?



# REVIEW

- ▶ How do we calculate linear regression coefficients?
- ▶ Minimize the sum of squared residuals.
- ▶ This is why linear regression is called Ordinary Least Squares (OLS).

$$\sum_{i=1}^n (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$



# REVIEW

$$\text{Residual}_i = \text{Observed}_i - \text{Prediction}_i$$



# REVIEW

$$\text{Residual}_i = y_i - \text{Prediction}_i$$



# REVIEW

$$\text{Residual}_i = y_i - (\widehat{\beta}_0 + \widehat{\beta}_1 x_i)$$



# REVIEW

$$\text{Residual}_i = y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i$$



# REVIEW

$$\text{Squared residual}_i = (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$



# REVIEW

$$\text{Sum of squared residuals} = \sum_{i=1}^n (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$





# REVIEW

- What do I need to know to calculate the sum of squared residuals?

$$\text{Sum of squared residuals} = \sum_{i=1}^n (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$

# REVIEW

- ▶ What do I need to know to calculate the sum of squared residuals?
- ▶ For every possible value of the coefficients, there is a single sum of squared residuals.

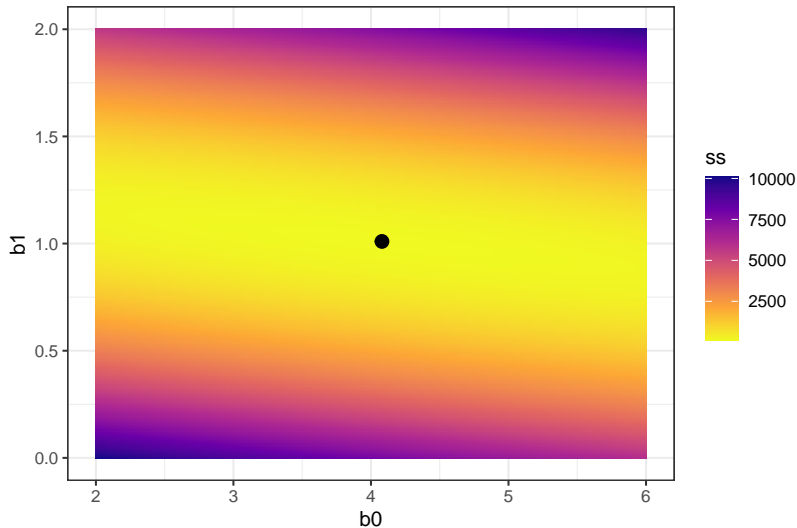
$$\text{Sum of squared residuals} = \sum_{i=1}^n (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$

# REVIEW

```
beta_grid <- as.data.table(expand.grid(b0=seq(2,6,0.01),b1=seq(0,2,0.01)))
ss <- function(i) {
  out <- data.table(b0=as.numeric(beta_grid[i,1]),
                    b1=as.numeric(beta_grid[i,2]),
                    ss=census[, sum((pcthouse-
                                     as.numeric(beta_grid[i,1])-
                                     as.numeric(beta_grid[i,2])*pctpop)^2)])
  return(out)
}
out <- rbindlist(lapply(1:nrow(beta_grid), ss))
```



# REVIEW



# REVIEW

```
out %>%  
  filter(ss==min(ss))
```

```
## Index: <ss>  
##      b0      b1      ss  
##    <num> <num>    <num>  
## 1:   4.08   1.01 144.8543
```

# REVIEW

```
## Outcome
Y <- as.matrix(census$pcthouse)
## Design matrix
X <- cbind(1, census$pcpop)
## Matrix algebra
solve(t(X) %*% X) %*% t(X) %*% Y
```

```
##           [,1]
## [1,] 4.081249
## [2,] 1.010295
```

# REVIEW

```
model <- lm(pcthouse~pctpop,  
            data=census)  
summary(model)
```

```
##  
## Call:  
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##  
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##      Min       1Q   Median       3Q      Max   
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## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
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How do we create a model based on empirical data?

$$pcthouse = f(pctpop)$$



How do we create a model based on empirical data?

$$pcthouse = f(pctpop)$$

1. Choose a functional form for the model.

$$pcthouse_i = \beta_0 + \beta_1 pctpop_i + \epsilon_i$$

# REVIEW

How do we create a model based on empirical data?

$$pcthouse = f(pctpop)$$

1. Choose a functional form for the model.

$$pcthouse_i = \beta_0 + \beta_1 pctpop_i + \epsilon_i$$

2. Choose an estimator (i.e., objective function) that relates the model to real observed data (e.g., OLS).

$$\text{Sum of squared residuals} = \sum_{i=1}^n (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$

# REVIEW

How do we create a model based on empirical data?

$$pcthouse = f(pctpop)$$

1. Choose a functional form for the model.

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$$\text{Sum of squared residuals} = \sum_{i=1}^n (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$

3. Fit the model (e.g., estimate parameters) by optimizing the objective function.

# REVIEW

How do we create a model based on empirical data?

$$pcthouse = f(pctpop)$$

1. Choose a functional form for the model.

$$pcthouse_i = \beta_0 + \beta_1 pctpop_i + \epsilon_i$$

2. Choose an estimator (i.e., objective function) that relates the model to real observed data (e.g., OLS).

$$\text{Sum of squared residuals} = \sum_{i=1}^n (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$

3. I want to maximize the likelihood that my observed data came from one particular model (e.g., set of parameters) relative to all other models.

57/73



## Why Language Models Hallucinate

Adam Tauman Kalai\*  
OpenAI

Ofir Nachum  
OpenAI

Santosh S. Vempala†  
Georgia Tech

Edwin Zhang  
OpenAI

September 4, 2025

### Abstract

Like students facing hard exam questions, large language models sometimes guess when uncertain, producing plausible yet incorrect statements instead of admitting uncertainty. Such “hallucinations” persist even in state-of-the-art systems and undermine trust. We argue that language models hallucinate because the training and evaluation procedures reward guessing over acknowledging uncertainty, and we analyze the statistical causes of hallucinations in the modern training pipeline. Hallucinations need not be mysterious—they originate simply as errors in binary classification. If incorrect statements cannot be distinguished from facts, then hallucinations in pretrained language models will arise through natural statistical pressures. We then argue that hallucinations persist due to the way most evaluations are graded—language models are optimized to be good test-takers, and guessing when uncertain improves test performance. This “epidemic” of penalizing uncertain responses can only be addressed through a socio-technical mitigation: modifying the scoring of existing benchmarks that are misaligned but dominate leaderboards, rather than introducing additional hallucination evaluations. This change may steer the field toward more trustworthy AI systems.

# PROPERTIES OF A GOOD ESTIMATOR

- ▶ How do we choose an objective function?



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- ▶ **Unbiased:** An unbiased estimator has an expected value equal to the “true” population parameter.



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- ▶ **Unbiased:** An unbiased estimator has an expected value equal to the “true” population parameter.
- ▶ **Consistency:** A consistent estimator “collapses” around the true value with variance zero as the sample size gets larger and moves towards infinity.





# PROPERTIES OF A GOOD ESTIMATOR

- ▶ How do we choose an objective function?
- ▶ **Unbiased:** An unbiased estimator has an expected value equal to the “true” population parameter.
- ▶ **Consistency:** A consistent estimator “collapses” around the true value with variance zero as the sample size gets larger and moves towards infinity.
- ▶ **Efficiency:** An efficient estimator is one that has a small sampling variance, relative to another estimator.



# PROPERTIES OF A GOOD ESTIMATOR

- ▶ There are entire classes on statistical inference that focus on deriving the proofs for these properties.



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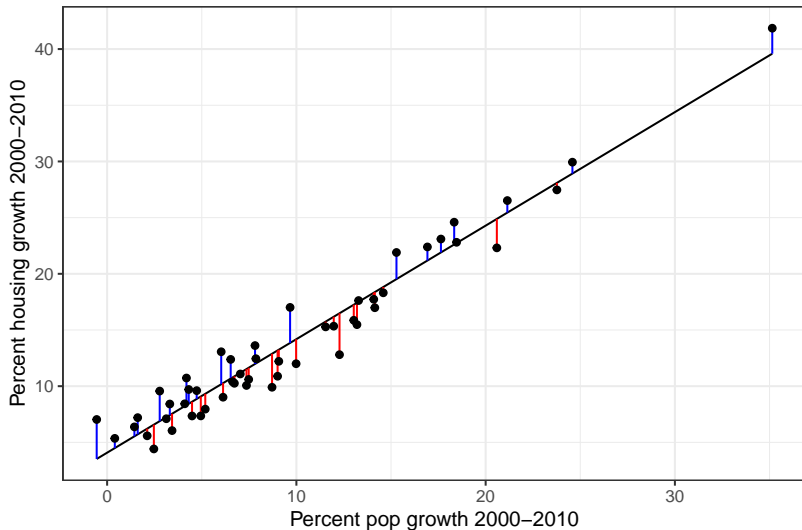


# PROPERTIES OF A GOOD ESTIMATOR

- ▶ There are entire classes on statistical inference that focus on deriving the proofs for these properties.
- ▶ The **Gauss-Markov theorem** states that for a linear regression model, the ordinary least squares (OLS) estimator provides the Best Linear Unbiased Estimator (BLUE), meaning it is the most precise (has the minimum variance) among all linear, unbiased estimators. This holds true when the model's error terms are uncorrelated, have equal variances, and a zero expectation, a set of assumptions known as **Gauss-Markov assumptions**.
- ▶ **We will discuss model diagnostics in Week 12.**

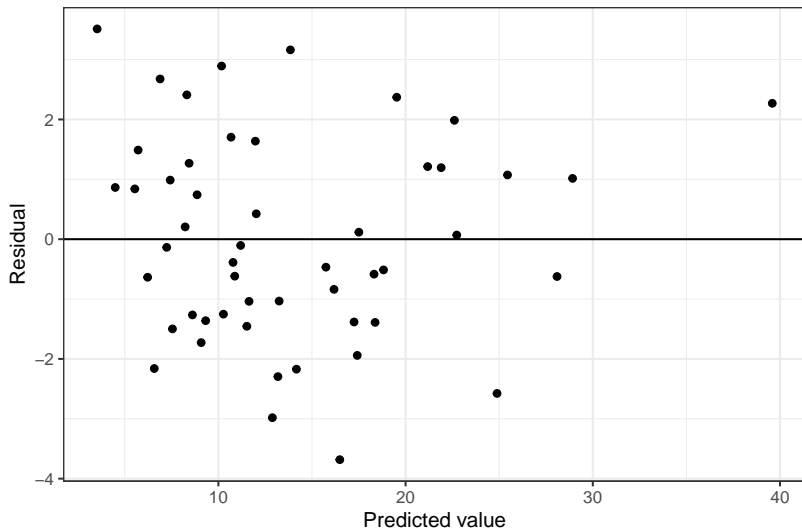


# RESIDUALS



# RESIDUALS

```
census <- census %>%  
  mutate(redisual=pcthouse-pcthouse_pred)  
  
plot <- ggplot(data=census,  
               aes(x=pcthouse_pred,  
                   y=residual)) +  
  
  geom_point() +  
  geom_hline(yintercept=0) +  
  labs(x='Predicted value', y='Residual') +  
  theme_bw()
```



# RESIDUALS

```
census %>%  
  summarize(mean_residual=round(mean(residual),4),  
            total_residual=round(sum(residual),4))
```

```
##      mean_residual total_residual  
## 1                0                0
```



# PROPERTIES OF OLS

- ▶ OLS produces residuals which are uncorrelated with predicted values.
- ▶ OLS produces residuals that sum to zero.



# GOODNESS OF FIT

```
summary(model)
```

```
##
## Call:
## lm(formula = pcthouse ~ pctpop, data = census)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.6830 -1.3132 -0.1364  1.2039  3.5126
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   4.08125    0.40793   10.01 1.98e-13 ***
## pctpop        1.01030    0.03371   29.97 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.719 on 49 degrees of freedom
## Multiple R-squared:  0.9483, Adjusted R-squared:  0.9472
## F-statistic: 898.1 on 1 and 49 DF,  p-value: < 2.2e-16
```

# INTERPRETING COEFFICIENTS

$$pcthouse_i = \beta_0 + \beta_1 pctpop_i + \epsilon_i$$

$$pcthouse_i = 4.08 + 1.01 pctpop_i + \epsilon_i$$

- How do we interpret the **intercept** coefficient?

# INTERPRETING COEFFICIENTS

$$pcthouse_i = \beta_0 + \beta_1 pctpop_i + \epsilon_i$$

$$pcthouse_i = 4.08 + 1.01 pctpop_i + \epsilon_i$$

- ▶ How do we interpret the **intercept** coefficient?
- ▶ How do we interpret the **slope** coefficient?

# INTERPRETING COEFFICIENTS

What if I divide my independent variable by 10?

```
census <- census %>%  
  mutate(pctpop_10 = pctpop/10)
```

# INTERPRETING COEFFICIENTS

What if I divide my independent variable by 10?

```
lm(pcthouse~pctpop_10, data=census)
```

```
##
```

```
## Call:
```

```
## lm(formula = pcthouse ~ pctpop_10, data = census
```

```
##
```

```
## Coefficients:
```

```
## (Intercept)      pctpop_10
```

```
##          4.081          10.103
```

# INTERPRETING COEFFICIENTS

What if I shift my independent variable by 10?

```
census <- census %>%  
  mutate(pctpop_10 = pctpop+10)  
lm(pcthouse~pctpop_10, data=census)
```

```
##
```

```
## Call:
```

```
## lm(formula = pcthouse ~ pctpop_10, data = census
```

```
##
```

```
## Coefficients:
```

```
## (Intercept)      pctpop_10
```

```
##      -6.022           1.010
```

# LINEAR TRANSFORMATIONS OF THE INDEPENDENT VARIABLE

## ► Rescaling



# LINEAR TRANSFORMATIONS OF THE INDEPENDENT VARIABLE

- ▶ Rescaling
- ▶ Shifting

# LINEAR TRANSFORMATIONS OF THE INDEPENDENT VARIABLE

- ▶ **Rescaling**
- ▶ **Shifting**
- ▶ Rescaling and shifting are useful for **interpretation** of regression coefficients. For example, say my dependent variable is life expectancy and my independent variable is income. How would I interpret the slope coefficient?

# LINEAR TRANSFORMATIONS OF THE INDEPENDENT VARIABLE

- ▶ **Rescaling**
- ▶ **Shifting**
- ▶ Rescaling and shifting are useful for **interpretation** of regression coefficients. For example, say my dependent variable is life expectancy and my independent variable is income. How would I interpret the slope coefficient?
- ▶ As we will see, they don't fundamentally change properties of the regression model or statistical tests.

# LINEAR TRANSFORMATIONS OF THE INDEPENDENT VARIABLE

- ▶ R-squared is the same.
- ▶ p-values are the same.

```
##
## Call:
## lm(formula = pcthouse ~ pctpop_10, data = census)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.6830 -1.3132 -0.1364  1.2039  3.5126
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -6.02170    0.70859  -8.498 3.34e-11 ***
## pctpop_10     1.01030    0.03371  29.968 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.719 on 49 degrees of freedom
## Multiple R-squared:  0.9483, Adjusted R-squared:  0.9472
## F-statistic: 898.1 on 1 and 49 DF,  p-value: < 2.2e-16
```

# LINEAR TRANSFORMATIONS OF THE INDEPENDENT VARIABLE

```
census <- census %>%  
  mutate(pctpop_rescaled = scale(pctpop))  
summary(lm(pcthouse~pctpop_rescaled, data=census))
```

```
##  
## Call:  
## lm(formula = pcthouse ~ pctpop_rescaled, data = census)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max   
## -3.6830 -1.3132 -0.1364  1.2039  3.5126   
##  
## Coefficients:  
##              Estimate Std. Error t value Pr(>|t|)      
## (Intercept)    13.9499     0.2408   57.94  <2e-16 ***  
## pctpop_rescaled  7.2868     0.2432   29.97  <2e-16 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 1.719 on 49 degrees of freedom  
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