### 431 Class 12

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# Previously Seen in 431

# Today's R Packages

```
library(NHANES)
library(car) # for Box-Cox transformation methods
library(janitor)
library(knitr)
library(broom)
library(magrittr)
library(patchwork)
library(ggrepel)
library(tidyverse)
theme set(theme bw())
```

# nh3\_new data (n = 989, 17 variables)

```
set.seed(20200914)
nh3 new <- NHANES %>%
    filter(SurveyYr == "2011_12") %>%
    select(ID, SurveyYr, Age, Height, Weight, BMI, Pulse,
           SleepHrsNight, BPSysAve, BPDiaAve, Gender,
           PhysActive, SleepTrouble, Smoke100,
           Race1, HealthGen, Depressed) %>%
    rename(Subject = ID, SleepHours = SleepHrsNight,
           Sex = Gender, SBP = BPSysAve, DBP = BPDiaAve) %>%
    filter(Age > 20 & Age < 80) %>%
    drop na() %>%
    distinct() %>%
    slice sample(n = 1000) %>%
    clean names() %>%
    filter(dbp > 39) %>%
    mutate(subject = as.character(subject))
```

# Today's Data (nh4)

```
set.seed(431)

nh4 <- nh3_new %>%
   select(subject, sbp, dbp, age, smoke100, race1) %>%
   slice_sample(n = 800, replace = FALSE)
```

- Outcome (quantitative): sbp
- Quantitative predictors: dbp, age
- Binary predictor: smoke100 (Yes/No)
- 5-category predictor: race1 (White, Black, Hispanic, Mexican, Other)
- Identification code: subject

# Models we've seen for sbp

### Create and relevel two additional variables

# Model mod\_4: add race\_white as a predictor

## **New Material**

# mod\_5: Using three race/ethnicity categories

```
Call:
```

```
lm(formula = sbp ~ dbp + age + smoke100 + race_3cat, data = nl
```

#### Coefficients:

```
      (Intercept)
      dbp
      age

      47.8831
      0.7449
      0.3835

      smoke100Yes
      race_3catBlack
      race_3catOther

      2.5655
      4.7147
      1.2232
```

OK. What's happened here? - What are our three categories for race\_3cat? - Why do I only see two of them in the model?

# Prediction for subject 65867?

subject	sbp	dbp	age	smoke100	race1	race_3cat
65867	115	78	60	No	White	White

 The referent category here is White, because that's the one left out of the set of indicators in the model. (We have coefficients for the other two race\_3cat categories.)

From Model 5, our predicted sbp for subject 65867 will be:

$$47.883 + 0.745 \; dbp + 0.384 \; age + 2.566 \; (if \; smoke100 = Yes) + 4.715 \; (if \; race\_3cat = Black) + 1.223 \; (if \; race\_3cat = Other)$$

So for subject 65867, we'd predict:

$$47.883 + 0.745$$
 (78) + 0.384 (60) + 2.566 (0) + 4.715 (0) + 1.223 (0) = 129.03 mm Hg

### augment for mod\_5

nh4\_aug5 <- augment(mod\_5, data = nh4)</pre>

subject	sbp	dbp	age	smoke100	race_3cat	.fitted	.resid
65867	115	78	60	No	White	128.9984	-13.998435
70046	125	83	55	No	White	130.8056	-5.805603
64302	98	59	45	No	White	109.0921	-11.092071
69386	141	68	52	Yes	Other	122.2697	18.730255

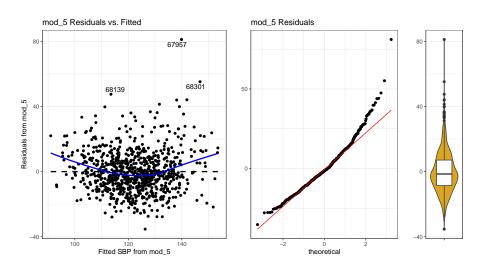
# Model mod\_5 results from tidy and glance

Coefficients for mod\_5 with 90% confidence intervals:

term	estimate	std.error	conf.low	conf.high
(Intercept)	47.88	3.20	42.61	53.15
dbp	0.74	0.04	0.68	0.81
age	0.38	0.03	0.33	0.43
smoke100Yes	2.57	0.93	1.03	4.10
race_3catBlack	4.71	1.33	2.53	6.90
race_3catOther	1.22	1.08	-0.55	3.00

r.squared	adj.r.squared	sigma	AIC	BIC
0.428	0.424	13	6378.7	6411.5

# Residual Plots for mod\_5?



# **Glancing at our Five Models**

model	preds	r.squared	adj.r.squared	sigma	AIC	BIC
1	dbp	0.291	0.290	14.40	6542.4	6556.4
2	1+age	0.414	0.413	13.10	6391.8	6410.6
3	2+smoke100	0.419	0.417	13.06	6387.3	6410.7
4	$3+$ race_white	0.424	0.421	13.01	6382.4	6410.5
5	3+race_3cat	0.428	0.424	12.97	6378.7	6411.5

Does there appear to be a clear winner here?

# Which one does best in our holdout sample?

We started with 989 subjects, and sampled 800 of them. How well do these models do when they are asked to predict the other 189 observations?

```
heldout <- anti_join(nh3_new, nh4, by = "subject") %>%
  select(subject, sbp, dbp, age, smoke100, race1) %>%
  mutate(race white = case when(race1 == "White" ~ 1.
                                     TRUE ~ ()) %>%
  mutate(race_3cat = fct_lump_n(race1, n = 2)) %>%
  mutate(race 3cat =
           fct relevel(race 3cat,
                       "White", "Black", "Other"))
dim(heldout)
```

[1] 189 8

## **Sanity Checks**

```
heldout %>% tabyl(race_white, race1)

race_white Black Hispanic Mexican White Other

0 38 18 17 0 15
1 0 0 0 101 0

heldout %>% tabyl(race_3cat, race1)
```

```
race_3cat Black Hispanic Mexican White Other
White 0 0 0 101 0
Black 38 0 0 0 0
Other 0 18 17 0 15
```

# How does our mod\_1 do out of sample?

```
heldout_mod1 <- augment(mod_1, newdata = heldout)
heldout_mod1 %>% select(subject, sbp, .fitted, .resid) %>%
  head() %>% kable()
```

subject	sbp	.fitted	.resid
65956	98	116.0260	-18.026024
71072	101	121.6898	-20.689797
64134	128	132.2082	-4.208233
66879	123	130.5900	-7.590012
66141	122	119.2625	2.737535
71279	147	150.0087	-3.008663

# Out-of-sample crude estimate of R-square

In our new sample, the square of the (Pearson) correlation between the observed sbp and the model mod\_1 predicted sbp or the .fitted values, will be our estimated R-square.

```
heldout_mod1 %$% cor(sbp, .fitted)
```

[1] 0.4841063

```
heldout_mod1 %$% cor(sbp, .fitted)^2
```

```
[1] 0.2343589
```

OK. So our estimate of the out-of-sample R-square = 0.234 based on this sample.

- How does this compare to our in-sample R-square for mod\_1, which was 0.291?
- Or maybe our adjusted R-square for mod\_1 which was 0.29?

## Create predictions for the other four models

```
heldout_mod2 <- augment(mod_2, newdata = heldout)
heldout_mod3 <- augment(mod_3, newdata = heldout)
heldout_mod4 <- augment(mod_4, newdata = heldout)
heldout_mod5 <- augment(mod_5, newdata = heldout)</pre>
```

# R<sup>2</sup> Comparisons for Models 1-5

Model	Predictors	In-sample R <sup>2</sup>	In-sample $R_{adj}^2$	Holdout R <sup>2</sup>
mod_1	dbp	0.291	0.29	0.234
mod_2	1 + age	0.414	0.413	0.329
mod_3	2 + smoke100	0.419	0.417	0.33
mod_4	$3 + {\tt race\_white}$	0.424	0.421	0.344
mod_5	$3 + race_3cat$	0.428	0.424	0.359

What if we look at the  $\sigma$  values - the residual standard deviations?

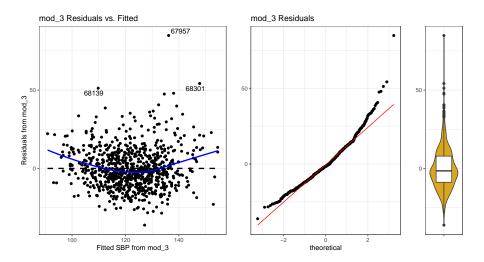
## $\sigma$ Comparisons for Models 1-5

Model	Predictors	In-sample $\sigma$	$Holdout\ \sigma$
mod_1	dbp	14.4	15.01
mod_2	1 + age	13.1	14.06
mod_3	2 + smoke100	13.06	14.04
mod_4	$3 + {\tt race\_white}$	13.01	13.9
mod_5	$3 + race_3cat$	12.97	13.73

Looks like our model summaries are just too optimistic?

• What might have tipped us off?

# Residual Plots (mod\_3)



# Why Transform the Outcome?

We want to try to identify a good transformation for the conditional distribution of the outcome, given the predictors, in an attempt to make the linear regression assumptions of linearity, Normality and constant variance more appropriate.

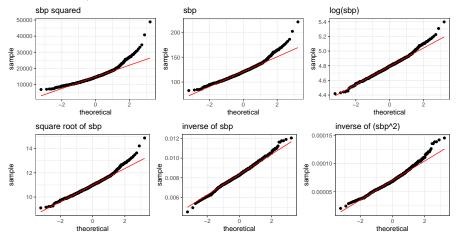
#### **Tukey's Ladder of Power Transformations**

Transformation	y <sup>2</sup>	У	$\sqrt{y}$	log(y)	$1/\sqrt{y}$	1/ <i>y</i>	$1/y^2$
$\lambda$	2	1	0.5	0	-0.5	-1	-2

• The most essential transformations (easy to understand) are the square, square root, logarithm and inverse.

## sbp distribution with various transformations





# Build model mod\_3 with log(sbp)

 Let's try a log transformation. We'll use the natural logarithm (log) as opposed to a base 10 logarithm (in R, log10) but that choice won't affect the residual plots.

```
mod_3log \leftarrow lm(log(sbp) \sim dbp + age + smoke100, data = nh4)
mod_3log
```

```
Call:
```

```
lm(formula = log(sbp) ~ dbp + age + smoke100, data = nh4)
```

#### Coefficients:

```
(Intercept) dbp age smoke100Yes
4.211704 0.006000 0.002989 0.019000
```

# Prediction for subject 65867?

subject	sbp	dbp	age	smoke100
65867	115	78	60	No

#### Call:

```
lm(formula = log(sbp) ~ dbp + age + smoke100, data = nh4)
```

#### Coefficients:

- Fitted log(sbp) = 4.21 + 0.006(78) + 0.003(60) + 0.019(0) = 4.859
- Observed log(sbp) = 4.745, so residual on the log scale is -0.104
- Predicted  $\mathbf{sbp} = \exp(4.859) = 128.9$ , while Observed  $\mathbf{sbp}$  was 115.

# Tidied coefficients of our log model

```
tidy(mod_3_log, conf.int = TRUE, conf.level = 0.90) %>%
  select(term, estimate, std.error, conf.low, conf.high) %>%
  kable(digits = 3)
```

term	estimate	std.error	conf.low	conf.high
(Intercept)	4.212	0.025	4.171	4.253
dbp	0.006	0.000	0.005	0.007
age	0.003	0.000	0.003	0.003
smoke100Yes	0.019	0.007	0.007	0.031

• Are these results comparable to our previous models?

# Fit summaries for our log model

```
glance(mod_3_log) %>%
  select(r.squared, adj.r.squared, sigma, AIC, BIC, nobs) %>%
  kable(digits = c(3,3,2,1,1,0))
```

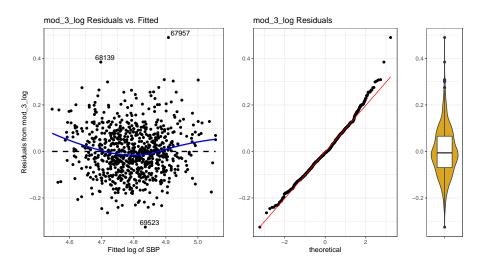
r.squared	adj.r.squared	sigma	AIC	BIC	nobs
0.428	0.425	0.1	-1367.6	-1344.1	800

• Are these results comparable to our previous models?

# Using augment with our logged model

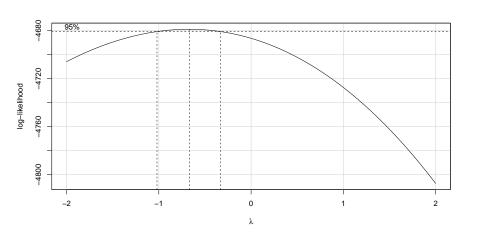
subject	sbp	log_sbp	.fitted	.resid	dbp	age	smoke100
65867	115	4.745	4.859	-0.114	78	60	No
70046	125	4.828	4.874	-0.046	83	55	No
64302	98	4.585	4.700	-0.115	59	45	No
69386	141	4.949	4.794	0.155	68	52	Yes

# Residual Plots for mod\_3\_log



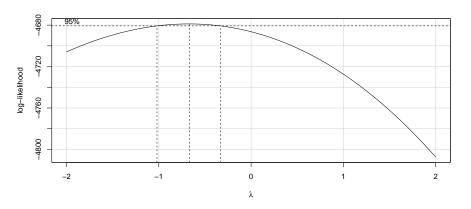
# Using Box-Cox to help select a transformation?

boxCox(mod\_3) # requires library(car)



# Remember the ladder!

Power $\lambda$	-2	-1	-0.5	0	0.5	1
transformation	$1/y^{2}$	1/y	$1/\sqrt{y}$	log(y)	$\sqrt{y}$	y



# Try the transformation suggested by Box-Cox?

Looks like  $1/\sqrt{sbp}$  (where  $\lambda=-0.5$ ) is the suggested transformation, although 1/sbp (where  $\lambda=-1$ ) is also within the reach of the provided 95% interval for  $\lambda$ .

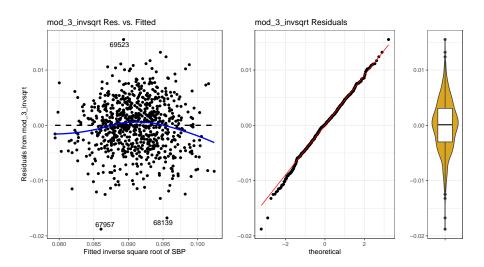
 As compared to the inverse square root, the inverse has the enormous advantage in many studies of being far easier to interpret.

Let's build each model and then we'll look at their residuals to see how well regression assumptions hold.

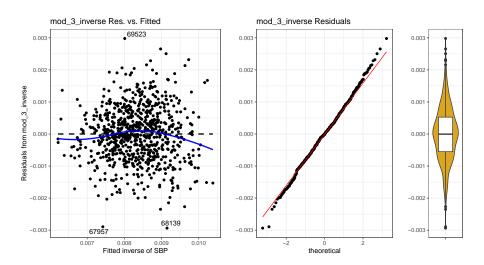
```
mod_3_invsqrt <-
lm((1/sqrt(sbp)) ~ dbp + age + smoke100, data = nh4)

mod_3_inverse <-
lm((1/sbp) ~ dbp + age + smoke100, data = nh4)</pre>
```

# Residual Plots for mod\_3\_invsqrt



## Residual Plots for mod\_3\_inverse



# **Early Notes on Transformations**

The Box-Cox plot (and indeed the ladder of power transformations) is designed to help identify transformations when:

- all of the values in the data are strictly positive, so that the log and square root, for instance, are defined
  - If your outcome includes zeros, you could just add 1 to each value before transforming at the risk of making interpretation harder.
- 2 the problems with assuming a linear relationship and/or Normality of residuals are indicated by more than just an outlier, or a few outliers.
- In that case, I would focus on the *influence* of those outliers (what happens to the model when you remove them?)

In any case, back-transforming predictions will be necessary at some stage if you apply a re-expression, which isn't too bad, but it can be challenging to write down the regression equation in terms of the original outcome.

### What have we discussed?

- The central role of linear regression in understanding associations between quantitative variables.
- The interpretation of a regression model as a prediction model.
- The meaning of key regression summaries, including residuals.
- Using tidy and glance from the broom package to help with summaries.
- Measuring association through correlation coefficients.
- How we might think about "adjusting" for the effect of a categorical predictor on a relationship between two quantitative ones.
- How a transformation might help us "linearize" the relationship shown in a scatterplot.