

# 432 Class 21 Slides

[github.com/THOMASELOVE/2019-432](https://github.com/THOMASELOVE/2019-432)

2019-04-16

# Preliminaries

```
library(skimr); library(MASS); library(robustbase)
library(quantreg); library(lmtest); library(sandwich)
library(boot); library(rms); library(survival)
library(OIsurv); library(survminer); library(broom)
library(tidyverse)
```

```
decim <- function(x, k) format(round(x, k), nsmall=k)
```

# Today's Agenda

- Regression on Time-to-event data
  - Cox Proportional Hazards Model
- Robust Linear Regression Methods
  - with Huber weights
  - with bisquare weights (biweights)
  - Bounded Influence Regression & Least Trimmed Squares
  - Penalized Least Squares using `ols` in `rms` package
  - Quantile Regression on the Median

# Survival Analysis / Cox Regression

# A Survival Analysis Example

Source: Chen and Peace (2011) *Clinical Trial Data Analysis Using R*, CRC Press, section 5.1

```
brca <- read.csv("data/breast_cancer.csv") %>% tbl_df
```

# The brca trial

The brca data describes a parallel randomized trial of three treatments, adjuvant to surgery in the treatment of patients with stage-2 carcinoma of the breast. The three treatment groups are:

- S+CT = Surgery plus one year of chemotherapy
- S+IT = Surgery plus one year of immunotherapy
- S+CT+IT = Surgery plus one year of chemotherapy and immunotherapy

The measure of efficacy were “time to death” in weeks. In addition to treat, our variables are:

- trial\_weeks: time in the study, in weeks, to death or censoring
- last\_alive: 1 if alive at last follow-up (and thus censored), 0 if dead
- age: age in years at the start of the trial

## brca tibble

```
# A tibble: 31 x 5
```

	subject	treat	trial_weeks	last_alive	age
	<fct>	<fct>	<int>	<int>	<int>
1	A01	S+CT	102	0	55
2	A02	S+IT	192	0	62
3	A03	S+CT+IT	73	0	72
4	A04	S+CT	58	1	48
5	A05	S+CT	48	1	26
6	A06	S+IT	182	1	52
7	A07	S+IT	196	1	50
8	A08	S+CT	177	1	49
9	A09	S+IT	191	1	62
10	A10	S+CT+IT	36	0	60

```
# ... with 21 more rows
```

# Analytic Objectives

This is a typical right-censored survival data set with interest in the comparative analysis of the three treatments.

- 1 Does immunotherapy added to surgery plus chemotherapy improve survival? (Comparing S+CT+IT to S+CT)
- 2 Does chemotherapy add efficacy to surgery plus immunotherapy? (S+CT+IT vs. S+IT)
- 3 What is the effect of age on survival?



# Create survival object

- `trial_weeks`: time in the study, in weeks, to death or censoring
- `last_alive`: 1 if alive at last follow-up (and thus censored), 0 if dead

So `last_alive = 0` if the event (death) occurs.

*What's next?*

# Create survival object

- `trial_weeks`: time in the study, in weeks, to death or censoring
- `last_alive`: 1 if alive at last follow-up (and thus censored), 0 if dead

So `last_alive = 0` if the event (death) occurs.

```
brca$S <- with(brca, Surv(trial_weeks, last_alive == 0))
```

```
head(brca$S)
```

```
[1] 102  192   73  58+  48+ 182+
```

# Build Kaplan-Meier Estimator

```
kmfit <- survfit(S ~ treat, dat = brca)
```

```
print(kmfit, print.rmean = TRUE)
```

```
Call: survfit(formula = S ~ treat, data = brca)
```

	n	events	*rmean	*se(rmean)	median	0.95LCL
treat=S+CT	11	6	153	21.1	144	102
treat=S+CT+IT	10	4	188	23.7	NA	139
treat=S+IT	10	5	188	17.9	192	144

0.95UCL

treat=S+CT	NA
treat=S+CT+IT	NA
treat=S+IT	NA

\* restricted mean with upper limit = 242

## summary(kmfit)

```
> summary(kmfit)
```

```
Call: survfit(formula = S ~ treat, data = brca)
```

treat=S+CT

time	n.risk	n.event	survival	std.err	lower 95% CI	upper 95% CI
55	10	1	0.900	0.0949	0.732	1.000
63	8	1	0.787	0.1340	0.564	1.000
102	7	1	0.675	0.1551	0.430	1.000
133	6	1	0.562	0.1651	0.316	1.000
144	5	1	0.450	0.1660	0.218	0.927
217	1	1	0.000	NaN	NA	NA

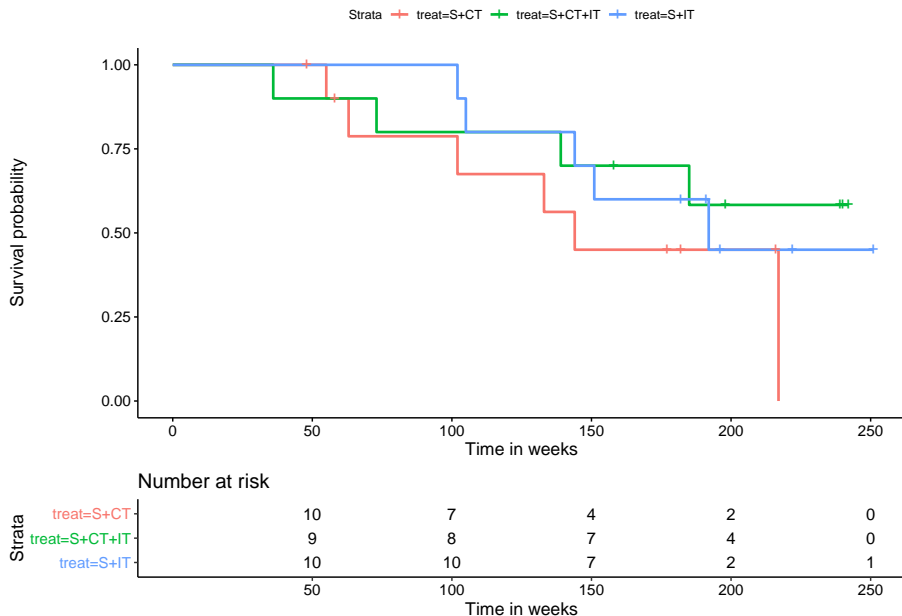
treat=S+CT+IT

time	n.risk	n.event	survival	std.err	lower 95% CI	upper 95% CI
36	10	1	0.900	0.0949	0.732	1
73	9	1	0.800	0.1265	0.587	1
139	8	1	0.700	0.1449	0.467	1
185	6	1	0.583	0.1610	0.340	1

treat=S+IT

time	n.risk	n.event	survival	std.err	lower 95% CI	upper 95% CI
102	10	1	0.90	0.0949	0.732	1.000
105	9	1	0.80	0.1265	0.587	1.000
144	8	1	0.70	0.1449	0.467	1.000

# K-M Plot via survminer



## K-M Plot via survminer (code)

```
ggsurvplot(kmfit, data = brca,  
            risk.table = TRUE,  
            risk.table.height = 0.25,  
            xlab = "Time in weeks")
```

# Testing the difference between curves

```
survdif(S ~ treat, dat = brca)
```

Call:

```
survdif(formula = S ~ treat, data = brca)
```

	N	Observed	Expected	$(O-E)^2/E$	$(O-E)^2/V$
treat=S+CT	11	6	3.80	1.2772	1.7647
treat=S+CT+IT	10	4	5.62	0.4676	0.7725
treat=S+IT	10	5	5.58	0.0605	0.0981

Chisq= 1.9 on 2 degrees of freedom, p= 0.4

What do we conclude?

## Fit Cox Model A: Treatment alone

```
modA <- coxph(S ~ treat, data = brca)
modA
```

Call:

```
coxph(formula = S ~ treat, data = brca)
```

	coef	exp(coef)	se(coef)	z	p
treatS+CT+IT	-0.8313	0.4355	0.6547	-1.270	0.204
treatS+IT	-0.5832	0.5581	0.6088	-0.958	0.338

Likelihood ratio test=1.75 on 2 df, p=0.4164  
n= 31, number of events= 15



## summary(modA)

```
> summary(modA)
```

Call:

```
coxph(formula = S ~ treat, data = brca)
```

n= 31, number of events= 15

	coef	exp(coef)	se(coef)	z	Pr(> z )
treatS+CT+IT	-0.8313	0.4355	0.6547	-1.270	0.204
treatS+IT	-0.5832	0.5581	0.6088	-0.958	0.338

	exp(coef)	exp(-coef)	lower .95	upper .95
treatS+CT+IT	0.4355	2.296	0.1207	1.571
treatS+IT	0.5581	1.792	0.1692	1.840

Concordance= 0.577 (se = 0.078 )

Rsquare= 0.055 (max possible= 0.944 )

Likelihood ratio test= 1.75 on 2 df, p=0.4164

Wald test = 1.82 on 2 df, p=0.403

Score (logrank) test = 1.89 on 2 df, p=0.3878

# Check Proportional Hazards Assumption

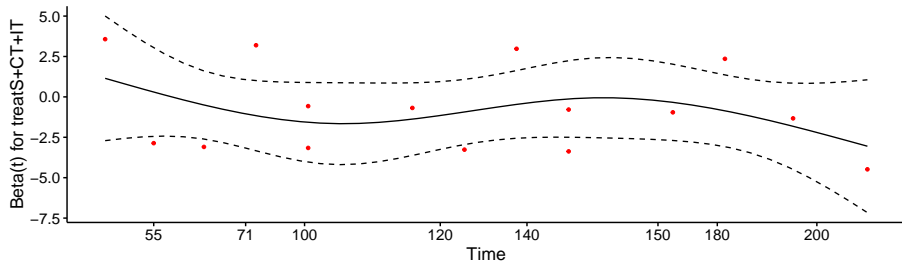
```
cox.zph(modA)
```

	rho	chisq	p
treatS+CT+IT	-0.198	0.618	0.432
treatS+IT	0.138	0.274	0.601
GLOBAL	NA	1.536	0.464

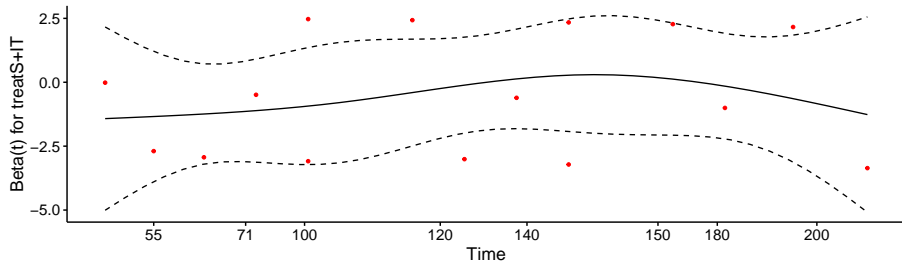
# Graphical PH Test `ggcoxzph(cox.zph(modA))`

Global Schoenfeld Test p: 0.4639

Schoenfeld Individual Test p: 0.4318



Schoenfeld Individual Test p: 0.6005



## Fit Cox Model B: Treatment + Age

```
modB <- coxph(S ~ treat + age, data = brca)
modB
```

Call:

```
coxph(formula = S ~ treat + age, data = brca)
```

	coef	exp(coef)	se(coef)	z	p
treatS+CT+IT	-0.59960	0.54903	0.65741	-0.912	0.3617
treatS+IT	-0.31161	0.73227	0.60936	-0.511	0.6091
age	0.07807	1.08119	0.03672	2.126	0.0335

Likelihood ratio test=6.99 on 3 df, p=0.07224

n= 31, number of events= 15

## summary(modB)

```
> summary(modB)
Call:
coxph(formula = S ~ treat + age, data = brca)

    n= 31, number of events= 15

              coef exp(coef) se(coef)      z Pr(>|z|)
treatS+CT+IT -0.59960   0.54903  0.65741 -0.912   0.3617
treatS+IT     -0.31161   0.73227  0.60936 -0.511   0.6091
age           0.07807   1.08119  0.03672  2.126   0.0335 *
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

              exp(coef) exp(-coef) lower .95 upper .95
treatS+CT+IT    0.5490    1.8214    0.1514    1.992
treatS+IT       0.7323    1.3656    0.2218    2.417
age             1.0812    0.9249    1.0061    1.162

Concordance= 0.701  (se = 0.083 )
Rsquare= 0.202  (max possible= 0.944 )
Likelihood ratio test= 6.99  on 3 df,   p=0.07224
Wald test            = 5.85  on 3 df,   p=0.1192
Score (logrank) test = 6.15  on 3 df,   p=0.1012
```

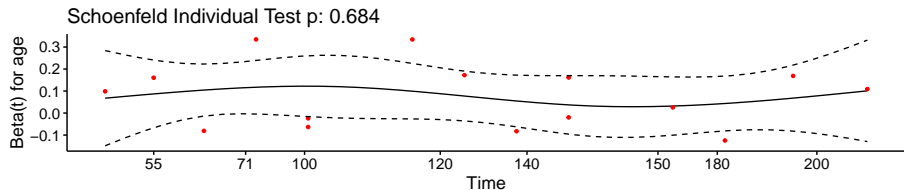
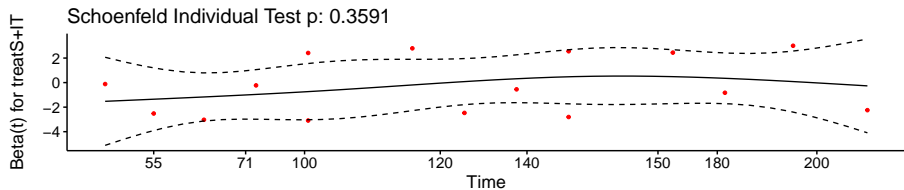
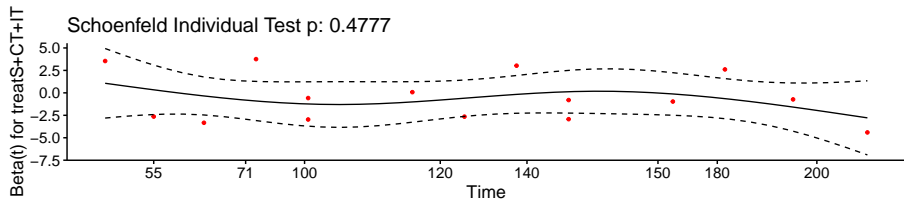
# Proportional Hazards Assumption: Model B Check

```
cox.zph(modB)
```

	rho	chisq	p
treatS+CT+IT	-0.179	0.504	0.478
treatS+IT	0.244	0.841	0.359
age	-0.106	0.166	0.684
GLOBAL	NA	2.416	0.491

# Graphical PH Test `ggcoxzph(cox.zph(modB))`

Global Schoenfeld Test p: 0.4907



# What to do if the PH assumption is violated

- If the PH assumption fails on a categorical predictor, fit a Cox model stratified by that predictor (use `strata(var)` rather than `var` in the specification of the `coxph` model.)
- If the PH assumption is violated, this means the hazard isn't constant over time, so we could fit separate Cox models for a series of time intervals.
- Use an extension of the Cox model that permits covariates to vary over time.

Visit

<https://cran.r-project.org/web/packages/survival/vignettes/timedep.pdf> for details on building the relevant data sets and models, with examples.



## The crimestat data and an OLS fit

# The crimestat data set

For each of 51 states (including the District of Columbia), we have the state's ID number, postal abbreviation and full name, as well as:

- **crime** - the violent crime rate per 100,000 people
- **poverty** - the official poverty rate (% of people living in poverty in the state/district) in 2014
- **single** - the percentage of households in the state/district led by a female householder with no spouse present and with her own children under 18 years living in the household in 2016
- **trump** - whether Donald Trump won the popular vote in the 2016 presidential election in that state/district (which we'll ignore for today)

# The crimestat data set

```
crimestat <- read.csv("data/crimestat.csv") %>% tbl_df(
  crimestat
```

```
# A tibble: 51 x 7
```

	sid	state	crime	poverty	single	trump	state.full
	<int>	<fct>	<dbl>	<dbl>	<dbl>	<int>	<fct>
1	1	AL	427.	19.2	9.02	1	Alabama
2	2	AK	636.	11.4	7.63	1	Alaska
3	3	AZ	400.	18.2	8.31	1	Arizona
4	4	AR	480.	18.7	9.41	1	Arkansas
5	5	CA	396.	16.4	7.25	0	California
6	6	CO	309.	12.1	6.75	0	Colorado
7	7	CT	237.	10.8	8.04	0	Connecticut
8	8	DE	489.	13	6.52	0	Delaware
9	9	DC	1244.	18.4	8.41	0	District of Colum~
10	10	FL	540.	16.6	8.29	1	Florida

```
# ... with 41 more rows
```

# Modeling crime with poverty and single

Our main goal will be to build a linear regression model to predict **crime** using centered versions of both **poverty** and **single**.

```
crimestat <- crimestat %>%  
  mutate(pov_c = poverty - mean(poverty),  
         single_c = single - mean(single))
```

# Our original (OLS) model

```
(mod1 <- lm(crime ~ pov_c + single_c, data = crimestat))
```

Call:

```
lm(formula = crime ~ pov_c + single_c, data = crimestat)
```

Coefficients:

(Intercept)	pov_c	single_c
364.41	16.11	23.84

# Significance of our coefficients?

```
tidy(mod1)
```

```
# A tibble: 3 x 5
```

	term	estimate	std.error	statistic	p.value
	<chr>	<dbl>	<dbl>	<dbl>	<dbl>
1	(Intercept)	364.	22.9	15.9	9.48e-21
2	pov_c	16.1	9.62	1.68	1.00e- 1
3	single_c	23.8	18.4	1.30	2.01e- 1

# Robust Linear Regression with Huber Weights

# Robust Linear Regression with Huber weights

There are several ways to do robust linear regression using M-estimation, including weighting using Huber and bisquare strategies.

- Robust linear regression here will make use of a method called iteratively re-weighted least squares (IRLS) to estimate models.
- M-estimation defines a weight function which is applied during estimation.
- The weights depend on the residuals and the residuals depend on the weights, so an iterative process is required.

We'll fit the model, using the default weighting choice: what are called Huber weights, where observations with small residuals get a weight of 1, and the larger the residual, the smaller the weight.

## Our robust model (using MASS::rlm)

```
rob.huber <- rlm(crime ~ pov_c + single_c, data = crimestat)
```



# Summary of the robust (Huber weights) model

```
tidy(rob.huber)
```

```
# A tibble: 3 x 4
```

	term	estimate	std.error	statistic
	<chr>	<dbl>	<dbl>	<dbl>
1	(Intercept)	344.	13.1	26.2
2	pov_c	11.9	5.51	2.16
3	single_c	31.0	10.5	2.94

Now, *both* predictors appear to have estimates that exceed twice their standard error. So this is a very different result than ordinary least squares gave us.

# Glance at the robust model (vs. OLS)

```
glance(mod1)
```

```
# A tibble: 1 x 11
  r.squared adj.r.squared sigma statistic p.value    df
    <dbl>         <dbl> <dbl>      <dbl>   <dbl> <int>
1    0.197         0.163  164.      5.88 0.00518     3
# ... with 5 more variables: logLik <dbl>, AIC <dbl>,
#   BIC <dbl>, deviance <dbl>, df.residual <int>
```

```
glance(rob.huber)
```

```
# A tibble: 1 x 6
  sigma converged logLik    AIC    BIC deviance
  <dbl> <lg1>      <dbl> <dbl> <dbl>   <dbl>
1  59.1 TRUE      -331.  671.  678. 1314784.
```

## Understanding the Huber weights a bit

Let's augment the data with results from this model, including the weights used.

```
crime_with_huber <- augment(rob.huber, crimestat) %>%  
  mutate(w = rob.huber$w) %>% arrange(w) %>% tbl_df  
  
head(crime_with_huber, 3)
```

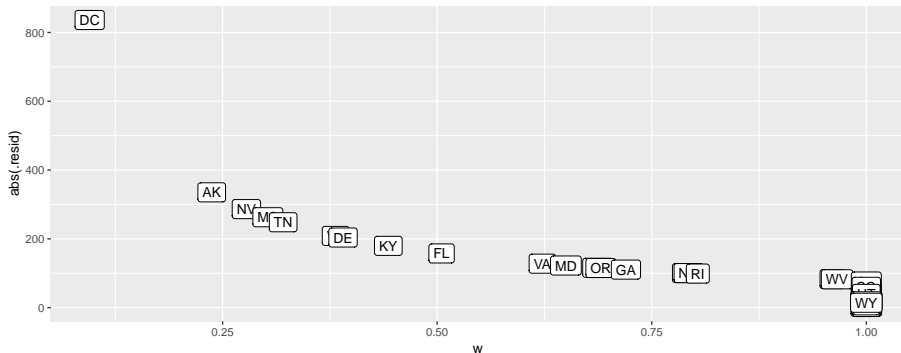
```
# A tibble: 3 x 15
```

	sid	state	crime	poverty	single	trump	state.full	pov_c
	<int>	<fct>	<dbl>	<dbl>	<dbl>	<int>	<fct>	<dbl>
1	9	DC	1244.	18.4	8.41	0	District ~	3.53
2	2	AK	636.	11.4	7.63	1	Alaska	-3.47
3	29	NV	636.	15.4	7.66	0	Nevada	0.527

```
# ... with 7 more variables: single_c <dbl>, .fitted <dbl>,  
#   .se.fit <dbl>, .resid <dbl>, .hat <dbl>, .sigma <dbl>,  
#   w <dbl>
```

# Are cases with large residuals down-weighted?

```
ggplot(crime_with_huber, aes(x = w, y = abs(.resid))) +  
  geom_label(aes(label = state))
```



# Conclusions from the Plot of Weights

- The district of Columbia will be down-weighted the most, followed by Alaska and then Nevada and Mississippi.
- But many of the observations will have a weight of 1.
- In ordinary least squares, all observations would have weight 1.
- So the more cases in the robust regression that have a weight close to one, the closer the results of the OLS and robust procedures will be.

## summary(rob.huber)

```
Call: rlm(formula = crime ~ pov_c + single_c, data = crimestat)
```

```
Residuals:
```

	Min	1Q	Median	3Q	Max
	-262.751	-45.641	1.762	36.732	836.244

```
Coefficients:
```

	Value	Std. Error	t value
(Intercept)	343.7982	13.1309	26.1823
pov_c	11.9098	5.5058	2.1631
single_c	30.9868	10.5266	2.9437

```
Residual standard error: 59.14 on 48 degrees of freedom
```

# Robust Linear Regression with the bisquare weighting function

## Robust Linear Regression with the biweight

As mentioned there are several possible weighting functions - we'll next try the biweight, also called the bisquare or Tukey's bisquare, in which all cases with a non-zero residual get down-weighted at least a little. Here is the resulting fit...

```
(rob.biweight <- rlm(crime ~ pov_c + single_c,  
                     data = crimestat, psi = psi.bisquare))
```

Call:

```
rlm(formula = crime ~ pov_c + single_c, data = crimestat, psi
```

Converged in 13 iterations

Coefficients:

(Intercept)	pov_c	single_c
336.17015	10.31578	34.70765

Degrees of freedom: 51 total; 48 residual

Scale estimate: 67.3



# Coefficients and Standard Errors

```
tidy(rob.biweight)
```

```
# A tibble: 3 x 4
```

	term	estimate	std.error	statistic
	<chr>	<dbl>	<dbl>	<dbl>
1	(Intercept)	336.	12.7	26.5
2	pov_c	10.3	5.31	1.94
3	single_c	34.7	10.2	3.42

# Understanding the biweights weights a bit

Let's augment the data, as above

```
crime_with_biweights <- augment(rob.biweight, crimestat) %>%  
  mutate(w = rob.biweight$w) %>% arrange(w) %>% tbl_df  
  
head(crime_with_biweights, 3)
```

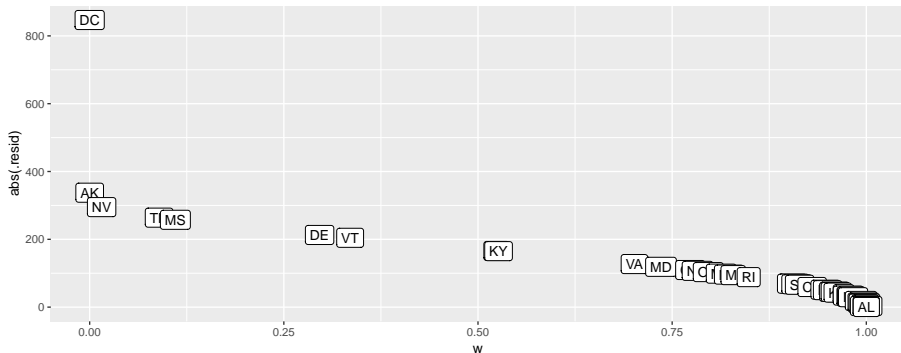
# A tibble: 3 x 13

	sid	state	crime	poverty	single	trump	state.full	pov_c
	<int>	<fct>	<dbl>	<dbl>	<dbl>	<int>	<fct>	<dbl>
1	2	AK	636.	11.4	7.63	1	Alaska	-3.47
2	9	DC	1244.	18.4	8.41	0	District ~	3.53
3	29	NV	636.	15.4	7.66	0	Nevada	0.527

# ... with 5 more variables: single\_c <dbl>, .fitted <dbl>,  
# .se.fit <dbl>, .resid <dbl>, w <dbl>

# Relationship of Weights and Residuals

```
ggplot(crime_with_biweights, aes(x = w, y = abs(.resid))) +  
  geom_label(aes(label = state))
```



# Conclusions from the biweights plot

Again, cases with large residuals (in absolute value) are down-weighted generally, but here, Alaska and Washington DC receive no weight at all in fitting the final model.

- We can see that the weight given to DC and Alaska is dramatically lower (in fact it is zero) using the bisquare weighting function than the Huber weighting function and the parameter estimates from these two different weighting methods differ.
- The maximum weight (here, for Alabama) for any state using the biweight is still slightly smaller than 1.

## summary(rob.biweight)

Call: `rlm(formula = crime ~ pov_c + single_c, data = crimestat`

Residuals:

Min	1Q	Median	3Q	Max
-257.58	-40.53	8.01	45.30	846.81

Coefficients:

	Value	Std. Error	t value
(Intercept)	336.1702	12.6733	26.5259
pov_c	10.3158	5.3139	1.9413
single_c	34.7077	10.1598	3.4162

Residual standard error: 67.27 on 48 degrees of freedom

# Comparing OLS and the two weighting schemes

```
glance(mod1) # OLS
```

```
# A tibble: 1 x 11
```

```
  r.squared adj.r.squared sigma statistic p.value    df  
    <dbl>      <dbl> <dbl>      <dbl>   <dbl> <int>  
1    0.197      0.163  164.      5.88 0.00518     3  
# ... with 5 more variables: logLik <dbl>, AIC <dbl>,  
#   BIC <dbl>, deviance <dbl>, df.residual <int>
```

```
glance(rob.biweight) # biweights
```

```
# A tibble: 1 x 6
```

```
  sigma converged logLik   AIC   BIC deviance  
  <dbl> <lgl>      <dbl> <dbl> <dbl>   <dbl>  
1  67.3 TRUE      -332.  672.  679. 1339850.
```

```
glance(rob.huber) # Huber weights
```

```
# A tibble: 1 x 6
```

# Bounded-Influence Regression

# Bounded-Influence Regression and Least-Trimmed Squares

Under certain circumstances, M-estimators can be vulnerable to high-leverage observations, and so, bounded-influence estimators, like least-trimmed squares (LTS) regression have been proposed. The biweight that we have discussed is often fitted as part of what is called an MM-estimation procedure, by using an LTS estimate as a starting point.

The `ltsReg` function, which is part of the `robustbase` package (Note: **not** the `ltsreg` function from MASS) is what I use below to fit a least-trimmed squares model. The LTS approach minimizes the sum of the  $h$  smallest squared residuals, where  $h$  is greater than  $n/2$ , and by default is taken to be  $(n + p + 1)/2$ .

## Least Trimmed Squares Model

```
lts1 <- ltsReg(crime ~ pov_c + single_c, data = crimestat)
```



# Summarizing the LTS model

```
summary(lts1)$coeff
```

	Estimate	Std. Error	t value	Pr(> t )
Intercept	339.14817	11.616766	29.194715	1.601245e-29
pov_c	16.99322	4.973459	3.416781	1.418337e-03
single_c	24.99819	9.136683	2.736024	9.073473e-03

# MM estimation

Specifying the argument `method="MM"` to `rlm` requests bisquare estimates with start values determined by a preliminary bounded-influence regression, as follows...

```
rob.MM <- rlm(crime ~ pov_c + single_c,  
              data = crimestat, method = "MM")
```

```
glance(rob.MM)
```

```
# A tibble: 1 x 6  
  sigma converged logLik    AIC    BIC deviance  
  <dbl> <lgl>      <dbl> <dbl> <dbl>    <dbl>  
1  75.8 TRUE      -332.  672.  679. 1337077.
```

## summary(rob.MM)

```
Call: rlm(formula = crime ~ pov_c + single_c, data = crimestat)
```

```
Residuals:
```

	Min	1Q	Median	3Q	Max
	-252.412	-41.096	8.696	47.141	847.128

```
Coefficients:
```

	Value	Std. Error	t value
(Intercept)	336.3928	13.1929	25.4980
pov_c	10.5579	5.5318	1.9086
single_c	32.7755	10.5763	3.0990

```
Residual standard error: 75.79 on 48 degrees of freedom
```

# Penalized Least Squares

# Penalized Least Squares with rms

We can apply a penalty to least squares directly through the `ols` function in the `rms` package.

```
d <- datadist(crimestat)
options(datadist = "d")
pls <- ols(crime ~ pov_c + single_c, penalty = 1,
           data = crimestat, x=T, y = T)
```

# The pls fit

## Linear Regression Model

```
ols(formula = crime ~ pov_c + single_c, data = crimestat, x =  
y = T, penalty = 1)
```

		Model Likelihood		Discrimination	
		Ratio Test		Indexes	
Obs	51	LR chi2	11.18	R2	0.197
sigma	159.1209	d.f.	1.946198	R2 adj	0.164
d.f.	48.0538	Pr(> chi2)	0.0035	g	89.298

## Residuals

Min	1Q	Median	3Q	Max
-284.24	-65.93	-16.68	15.66	807.01

# How to Choose the Penalty in Penalized Least Squares?

The problem here is how to choose the penalty - and that's a subject I'll essentially skip today. The most common approach (that we've seen with the lasso) is cross-validation.

Meanwhile, what do we conclude about the fit here from AIC and BIC?

```
AIC(pls); BIC(pls)
```

```
d.f.
```

```
669.5781
```

```
d.f.
```

```
677.2014
```

## Quantile Regression (on the Median)



# Quantile Regression on the Median

We can use the `rq` function in the `quantreg` package to model the **median** of our outcome (violent crime rate) on the basis of our predictors, rather than the mean, as is the case in ordinary least squares.

```
rob.quan <- rq(crime ~ pov_c + single_c, data = crimestat)

glance(rob.quan)
```

```
# A tibble: 1 x 5
  tau logLik   AIC   BIC df.residual
<dbl> <dbl> <dbl> <dbl>      <int>
1  0.5 -316.  638.  643.         48
```

## summary(rob.quan)

```
Call: rq(formula = crime ~ pov_c + single_c, data = crimestat)
```

```
tau: [1] 0.5
```

Coefficients:

	coefficients	lower bd	upper bd
(Intercept)	344.75658	336.94534	366.23603
pov_c	10.54757	3.06714	28.95962
single_c	32.27249	4.45889	48.18925

## Estimating a different quantile ( $\tau = 0.70$ )

In fact, if we like, we can estimate any quantile by specifying the  $\tau$  parameter (here  $\tau = 0.5$ , by default, so we estimate the median.)

```
(rob.quan70 <- rq(crime ~ pov_c + single_c, tau = 0.70,  
                  data = crimestat))
```

Call:

```
rq(formula = crime ~ pov_c + single_c, tau = 0.7, data = crimestat)
```

Coefficients:

(Intercept)	pov_c	single_c
379.72818	19.30376	32.15827

Degrees of freedom: 51 total; 48 residual

# Conclusions

# Comparing Five of the Models

## Estimating the Mean

	Fit	Intercept CI	pov_c CI	single_c CI
	OLS	(318.6, 410.2)	(-3.13, 35.35)	(-12.92, 60.60)
	Robust (Huber)	(320.0, 367.6)	(0.89, 22.93)	(9.93, 52.05)
	Robust (biweight)	(310.7, 361.5)	(-0.30, 20.94)	(14.39, 55.03)
	Robust (MM)	(310.0, 362.8)	(-0.50, 21.62)	(11.62, 53.94)

**Note:** CIs estimated for OLS and Robust methods as point estimate  $\pm$  2 standard errors

## Estimating the Median

	Fit	Intercept CI	pov_c CI	single_c CI
Quantile (Median) Reg		(336.9, 366.2)	(3.07, 28.96)	(4.46, 48.19)

# Comparing AIC and BIC

Fit	AIC	BIC
OLS	669.7	677.4
Robust (Huber)	670.8	678.5
Robust (biweight)	671.7	679.4
Robust (MM)	671.6	679.3
Quantile (median)	637.5	643.3

# Some General Thoughts

- 1 When comparing the results of a regular OLS regression and a robust regression for a data set which displays outliers, if the results are very different, you will most likely want to use the results from the robust regression.
  - Large differences suggest that the model parameters are being highly influenced by outliers.
- 2 Different weighting functions have advantages and drawbacks.
  - Huber weights can have difficulties with really severe outliers.
  - Bisquare weights can have difficulties converging or may yield multiple solutions.
  - Quantile regression approaches have some nice properties, but describe medians (or other quantiles) rather than means.