

# Inclass\_\_week10

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```
library(tidyverse)

## -- Attaching packages ----- tidyverse 1.2.1 --
## v ggplot2 3.1.0      v purrr  0.2.5
## v tibble  2.0.1      v dplyr  0.7.8
## v tidyr   0.8.2      v stringr 1.3.1
## v readr   1.3.1      v forcats 0.3.0

## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()

library(modelr)
library(caret)

## Loading required package: lattice

##
## Attaching package: 'caret'

## The following object is masked from 'package:purrr':
##
## lift

load("els.Rdata")

## Define the model
mod1_formula<-formula(bynels2m~byses1+ ##DEFINE YOUR OWN MODEL HERE
                      bynels2r)
## Run the model against all of the data
basic.mod<-lm(mod1_formula,
              data=els); summary(basic.mod)

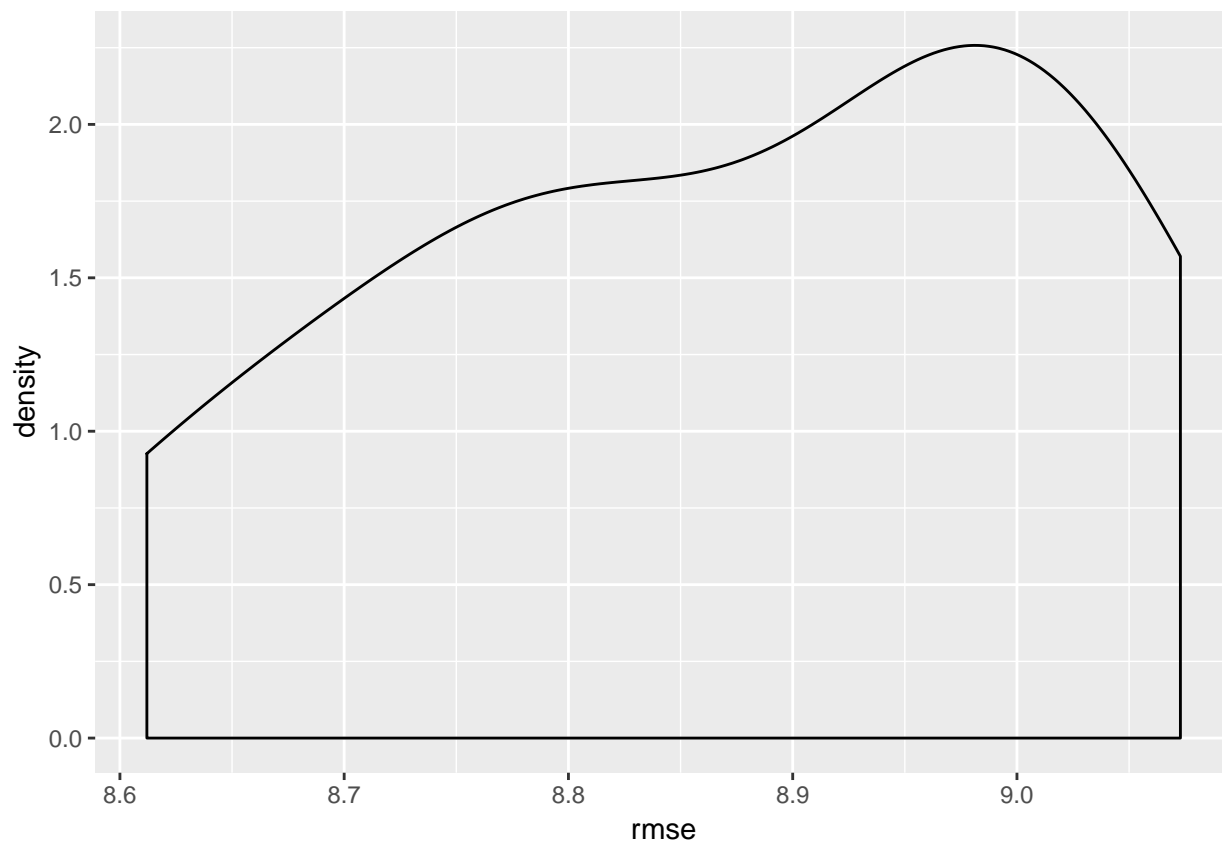
##
## Call:
## lm(formula = mod1_formula, data = els)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -32.212  -5.992  -0.165   5.853  46.062
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 15.844996   0.256915   61.67  <2e-16 ***
## byses1       2.493117   0.106930   23.32  <2e-16 ***
## bynels2r     0.991916   0.008382  118.34  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8.871 on 15322 degrees of freedom
```

```
## (964 observations deleted due to missingness)
## Multiple R-squared: 0.5759, Adjusted R-squared: 0.5758
## F-statistic: 1.04e+04 on 2 and 15322 DF, p-value: < 2.2e-16

els_cf<-els%>% ##SUBSTITUTE YOUR OWN DATA HERE
  crossv_kfold(10)

rmse_mod1<-els_cf %>%
  mutate(train = map(train, as_tibble)) %>% ## Convert to tibbles
  mutate(model = map(train, ~ lm(mod1_formula,
                                data = .))) %>%
  mutate(rmse = map2_dbl(model, test, rmse)) %>% ## apply model, get rmse
  select(.id, rmse) ## pull just id and rmse

gg<-ggplot(data=rmse_mod1,aes(x=rmse))
gg<-gg+geom_density()
gg
```



```
els_cv<-els%>% ##SUBSTITUTE YOUR OWN DATA HERE
  crossv_mc(n=1000,test=.2)
els_cv
```

```
## # A tibble: 1,000 x 3
##   train      test      .id
##   <list>     <list>   <chr>
## 1 <S3: resample> <S3: resample> 0001
## 2 <S3: resample> <S3: resample> 0002
## 3 <S3: resample> <S3: resample> 0003
## 4 <S3: resample> <S3: resample> 0004
```

```
## 5 <S3: resample> <S3: resample> 0005
## 6 <S3: resample> <S3: resample> 0006
## 7 <S3: resample> <S3: resample> 0007
## 8 <S3: resample> <S3: resample> 0008
## 9 <S3: resample> <S3: resample> 0009
## 10 <S3: resample> <S3: resample> 0010
## # ... with 990 more rows
```

```
mod1_rmse_cv<-els_cv %>%
  mutate(train = map(train, as_tibble)) %>% ## Convert to tibbles
  mutate(model = map(train, ~ lm(mod1_formula, data = .)))%>%
  mutate(rmse = map2_dbl(model, test, rmse))%>%
  select(.id, rmse) ## pull just id and rmse
```

```
summary(mod1_rmse_cv$rmse)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      8.496   8.797   8.880   8.877   8.953   9.311
```

```
gg<-ggplot(mod1_rmse_cv,aes(x=rmse))
gg<-gg+geom_density()
gg
```

