Sums of Squares in R

uoepsy.github.io

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After a bit of back and forth in the Live R session about which functions in R give you which type of sums of squares (and indeed us misremembering which type is which!), we figured we (and hopefully you too) would benefit from a small explainer.

Some Data

```
library(tidyverse)
usmrsurv2 <- read_csv("https://uoepsy.github.io/data/usmrsurvey2.csv")</pre>
names(usmrsurv2)
    [1] "id"
                                "pseudonym"
                                                       "catdog"
                                                       "optimism"
    [4] "gender"
                                "height"
    [7] "spirituality"
                               "ampm"
                                                       "extraversion"
## [10] "agreeableness"
                               "conscientiousness"
                                                       "emotional_stability"
                               "internal_control"
## [13] "imagination"
names(usmrsurv2)[9:14]<-c("E","A","C","ES","I","LOC")
```

A model

Here is a model with two predictors, with the order of the predictors differing between the two models:

```
mymod1 <- lm(LOC ~ ES + optimism, data = usmrsurv2)
mymod2 <- lm(LOC ~ optimism + ES, data = usmrsurv2)</pre>
```

Types of Sums of Squares

Type 1

Type 1 Sums of Squares is the "incremental" or "sequential" sums of squares. If we have a model $Y \sim A + B$, this method tests:

- the main effect of A
- the main effect of B after the main effect of A
- Interactions (which come in Week 9 of the course) are tested after the main effects

Because this is sequential, the order matters.

We can get the Type 1 SS in R using the function anova().

As you will see, the order in which the predictors are entered in the model influences the results. This is because:

- for mymod1 (lm(LOC ~ ES + optimism)) we are testing the main effect of ES, followed by the main effect of optimism after accounting for effects of ES.
- for mymod2 (lm(LOC ~ optimism + ES)) it is the other way around: we test the main effect of optimism, followed by the main effect of ES after accounting for effects of optimism.

```
# mymod1 <- lm(LOC ~ ES + optimism, data = usmrsurv2)</pre>
anova(mymod1)
## Analysis of Variance Table
##
## Response: LOC
            Df Sum Sq Mean Sq F value
##
             1 114.66 114.661 7.3357 0.009036 **
             1 16.80 16.797 1.0746 0.304527
## optimism
## Residuals 54 844.05 15.631
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
# mymod2 <- lm(LOC ~ optimism + ES, data = usmrsurv2)</pre>
anova(mymod2)
## Analysis of Variance Table
##
## Response: LOC
##
            Df Sum Sq Mean Sq F value Pr(>F)
## optimism
            1 31.81 31.813 2.0353 0.15944
             1 99.64 99.644 6.3749 0.01454 *
## Residuals 54 844.05 15.631
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Type 3

Type 3 Sums of Squares is the "partial" sums of squares. If we have a model $Y \sim A + B$, this method tests:

- the main effect of A after the main effect of B
- the main effect of B after the main effect of A

So the Type 3 will be equivalent to Type 1 only for the final predictor in the model.

Martin showed us one approach in the Live R session, using the drop1() function:

```
# mymod1 <- lm(LOC ~ ES + optimism, data = usmrsurv2)</pre>
drop1(mymod1, test = "F")
## Single term deletions
##
## Model:
## LOC ~ ES + optimism
           Df Sum of Sq
##
                            RSS
                                  AIC F value Pr(>F)
## <none>
                         844.05 159.62
                  99.644 943.70 163.99 6.3749 0.01454 *
## ES
             1
                  16.797 860.85 158.75 1.0746 0.30453
## optimism 1
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Note that these results are the same as the Type 3 for optimism in the model $lm(LOC \sim ES + optimism)$, and the Type 3 for ES in the model $lm(LOC \sim optimism + ES)$. Take a look at the previous page for confirmation of this.

Note that Type 3 SS are **invariant to the order of predictors:** We get the same when we switch around our predictors:

```
# mymod2 <- lm(LOC ~ optimism + ES, data = usmrsurv2)</pre>
drop1(mymod2, test = "F")
## Single term deletions
##
## Model:
## LOC ~ optimism + ES
##
           Df Sum of Sq
                           RSS
                                   AIC F value Pr(>F)
                         844.05 159.62
## <none>
## optimism 1
                  16.797 860.85 158.75 1.0746 0.30453
## ES
                  99.644 943.70 163.99 6.3749 0.01454 *
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

The summary() function

Remember that we mentioned in the Week 8 lab that in the simple regression model (one predictor), the t-statistic for the coefficient test is the square root of the F-statistic for the test of the overall reduction in the residual sums of squares?

Well this does still hold for the multiple regression model, but it is a little more complicated.

For a given coefficient t-statistic in a multiple regression model, the associated F-statistic is the one corresponding to the reduction in residual sums of squares that is attributable to that predictor only. Or, in other words, the Type 3 F-statistic.

Here are our model coefficients and t-statistics:

```
summary(mymod1)$coefficients
```

```
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 15.7589278 2.32636156 6.774066 9.587579e-09
## ES 0.1904987 0.07544910 2.524863 1.454432e-02
## optimism 0.0223095 0.02152114 1.036632 3.045273e-01
```

Here are our Type 3 SS F-statistics:

```
drop1(mymod1, test = "F")
```

```
## Single term deletions
##
## Model:
## LOC ~ ES + optimism
##
           Df Sum of Sq
                           RSS
                                  AIC F value Pr(>F)
## <none>
                        844.05 159.62
## ES
                 99.644 943.70 163.99
                                       6.3749 0.01454 *
                 16.797 860.85 158.75 1.0746 0.30453
## optimism
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

We can square-root them to get back to the t-statistic:

```
sqrt(drop1(mymod1, test = "F")$`F value`)
```

```
## [1] NA 2.524863 1.036632
```

What this means is that just like the drop1() F test for reduction in residual sums of squares uses Type 3 SS, the t tests for the coefficients produced in summary() for a linear model are also invariant to the order of predictors.