



# ANCOVA

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# Inhoud

- ancova 1
  - define the model
  - group means
  - model fit no covar
  - model fit only covar
  - model fit with covar
  - total variance
  - total variance visual
  - model variance group visual
  - model variance covariate visual
  - model variance group and covariate visual
  - error variance with covariate
  - sums of squares
  - f ratio
  - adjusted means



# ANCOVA

# ANCOVA

Analysis of covariance (ANCOVA) is a general linear model which blends ANOVA and regression. ANCOVA evaluates whether population means of a dependent variable (DV) are equal across levels of a categorical independent variable (IV) often called a treatment, while statistically controlling for the effects of other continuous variables that are not of primary interest, known as covariates (CV).

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# ANCOVA

Determine main effect while correcting for covariate

- 1 dependent variable
- 1 or more independent variables
- 1 or more covariates

A covariate is a variable can be a confounding variable biasing your results. By adding a covariate we reduce error/residual in the model.



# Assumptions

- Same as ANOVA
- Independence of the covariate and treatment effect §12.3.1.
  - No difference on ANOVA with covar and independent variable
  - Matching experimental groups on the covariate
- Homogeneity of regression slopes §12.3.2.
  - Visual: scatterplot dep var \* covar per condition
  - Testing: interaction indep. var \* covar



# Data example

We want to test the difference in national extraversion but want to control for openness to experience.

- Dependent variable: Extraversion
- Independent variable: Nationality
  - Dutch
  - German
  - Belgian
- Covariate: Openness to experience



# Simulate data

```
# Simulate data
n = 20
k = 3
group      = round(runif(n,1,k),0)
mu.covar   = 8
sigma.covar = 1
covar       = round(rnorm(n,mu.covar,sigma.covar),2)

# Create dummy variables
dummy.1 <- ifelse(group == 2, 1, 0)
dummy.2 <- ifelse(group == 3, 1, 0)

# Set parameters
b.0 = 15 # initial value for group 1
b.1 = 3  # difference between group 1 and 2
b.2 = 4  # difference between group 1 and 3
b.3 = 3  # Weight for covariate

# Create error
error = rnorm(n,0,1)
```



# Define the model

$outcome = model + error$

$model = indvar + covariate = nationality + openness$

Formal model

$b_0 + b_1 dummy_1 + b_2 dummy_2 + b_3 covar$

```
# Define model  
outcome = b.0 + b.1 * dummy.1 + b.2 * dummy.2 + b.3 * covar + error
```



# Dummies



# The data



# Group means

```
aggregate(outcome ~ group, data, mean)
```

```
##   group  outcome
## 1      1 38.67000
## 2      2 42.16625
## 3      3 43.79800
```



# Model fit no covar

What are the beta coëfficients based on the data without the covariate?

```
fit.group <- lm(outcome ~ factor(group), data); fit.group
```

```
##  
## Call:  
## lm(formula = outcome ~ factor(group), data = data)  
##  
## Coefficients:  
## (Intercept) factor(group)2 factor(group)3  
## 38.670       3.496       5.128
```

```
fit.group$coefficients[2:3] + fit.group$coefficients[1]
```

```
## factor(group)2 factor(group)3  
## 42.16625      43.79800
```



# Model fit only covar

What is the weight of only the covariate?

```
fit.covar <- lm(outcome ~ covar, data)
fit.covar
```

```
##
## Call:
## lm(formula = outcome ~ covar, data = data)
##
## Coefficients:
## (Intercept)      covar
##       15.667        3.185
```



# Model fit with covar

```
fit <- lm(outcome ~ factor(group) + covar, data); fit
```

```
##  
## Call:  
## lm(formula = outcome ~ factor(group) + covar, data = data)  
##  
## Coefficients:  
## (Intercept) factor(group)2 factor(group)3 covar  
## 15.965 2.769 4.181 2.881
```

```
fit$coefficients[2:3] + fit$coefficients[1]
```

```
## factor(group)2 factor(group)3  
## 18.73401 20.14609
```

$$\text{Dutch} = 15.96 \text{ German} = 18.73 \text{ Belgian} = 20.14$$

# Total variance

What is the total variance

$$MS_{total} = s_{outcome}^2 = \frac{SS_{outcome}}{df_{outcome}}$$

```
ms.t = var(data$outcome); ms.t
```

```
## [1] 11.97756
```

```
ss.t = var(data$outcome) * (length(data$outcome) - 1); ss.t
```

```
## [1] 227.5737
```



# The data

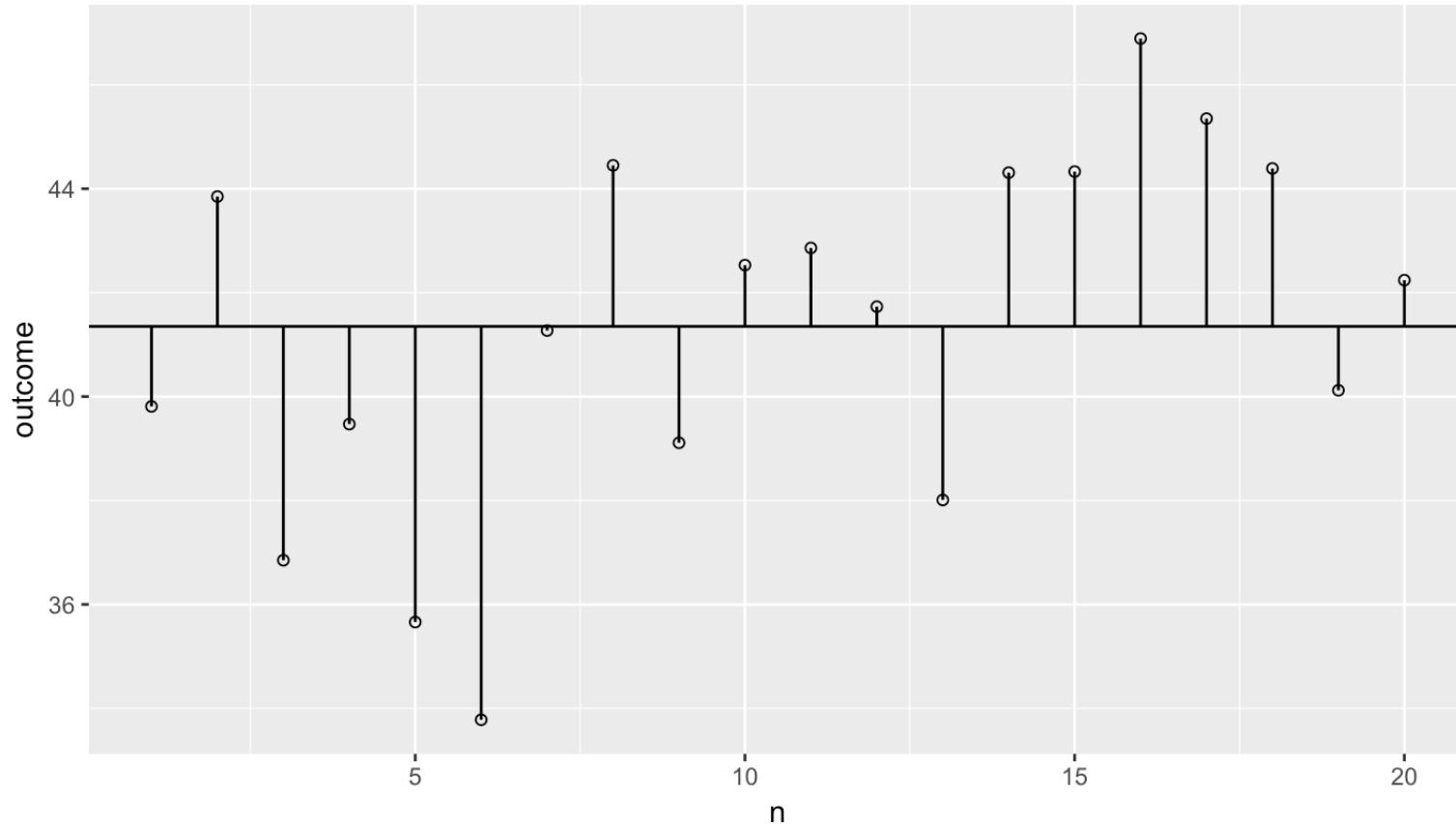


# Total variance visual

```
plot <- ggplot(data, aes(x=n, y=outcome)) + geom_point(shape=1) +  
  geom_hline(yintercept=mean(outcome)) +  
  geom_segment(aes(x = n, y = grand.mean, xend = n, yend = outcome)) +  
  ggtitle("Total variance")  
plot
```



## Total variance



# Model variance group

The model variance consists of two parts. One for the independent variable and one for the covariate. Lets first look at the independent variable.

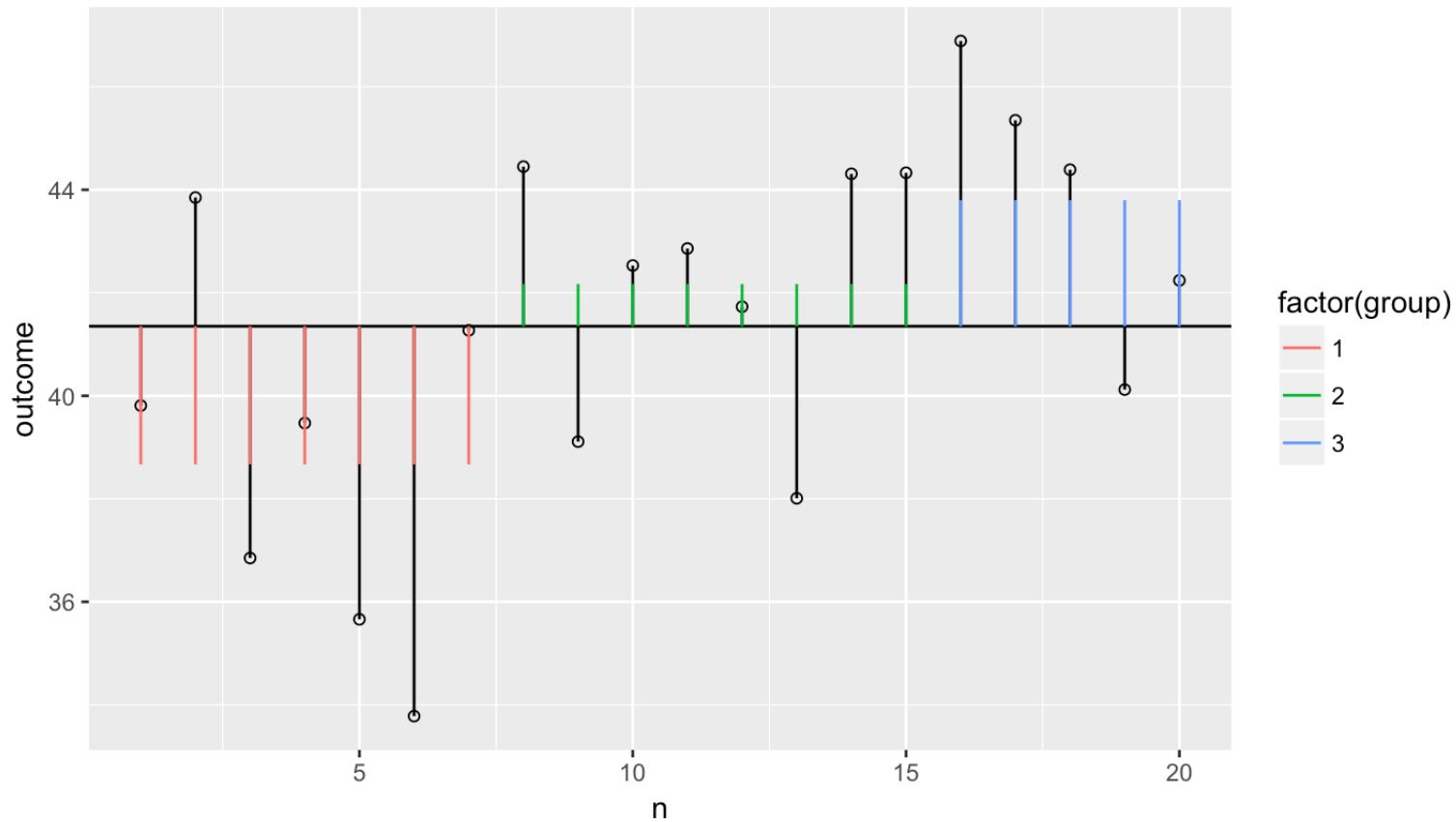


# Model variance group visual

```
plot + geom_segment(aes(x = n, y = grand.mean, xend = n, yend = data$model.gro  
gtitle("Only group variance")
```



## Only group variance

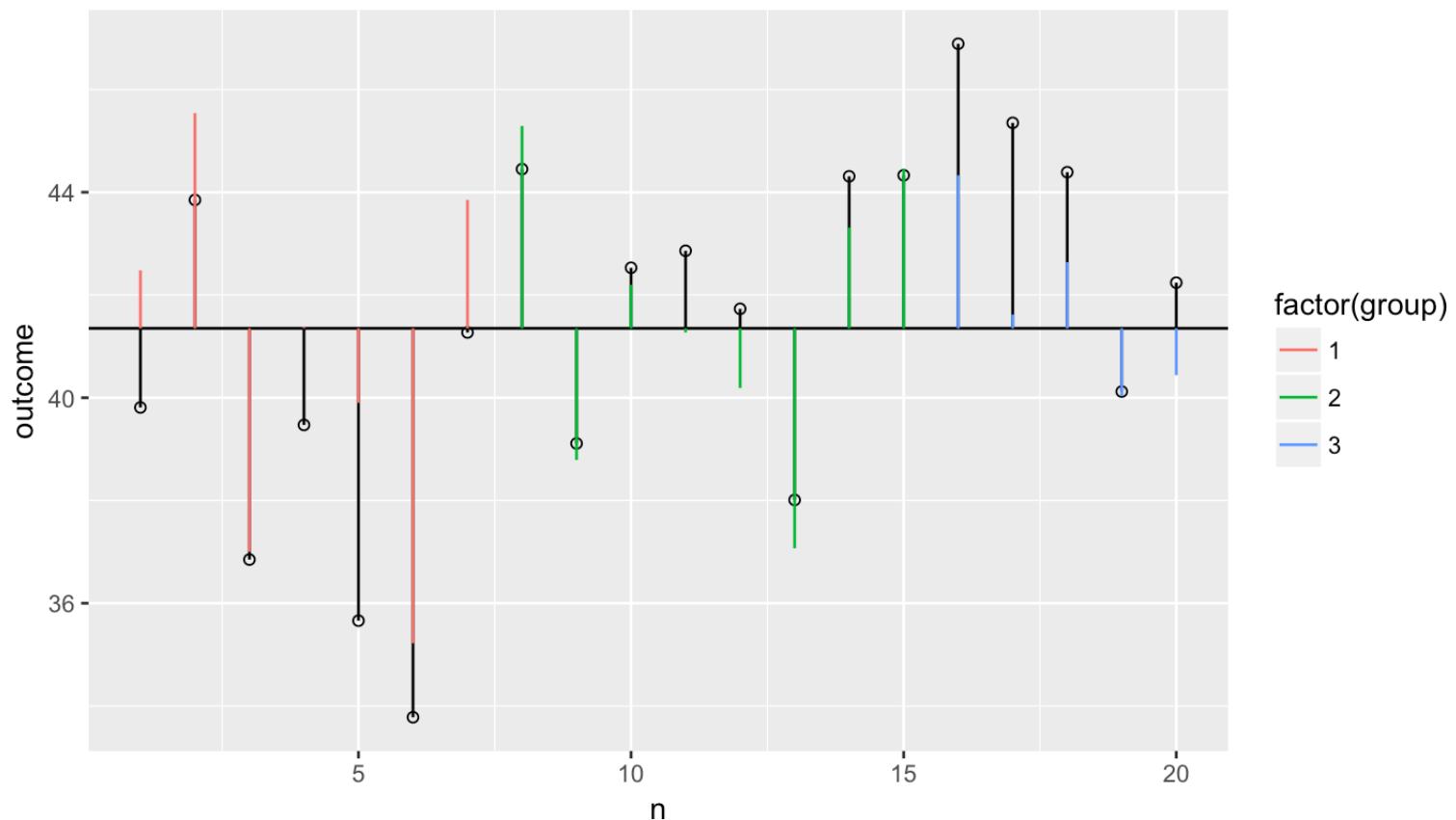


# Model variance covariate visual

```
data$model.covar <- round(fit.covar$fitted.values,2)  
plot + geom_segment(aes(x = n, y = grand.mean, xend = n, yend = data$model.cov  
ggtile("Only covariate variance")
```



## Only covariate variance



# Model variance group and covariate

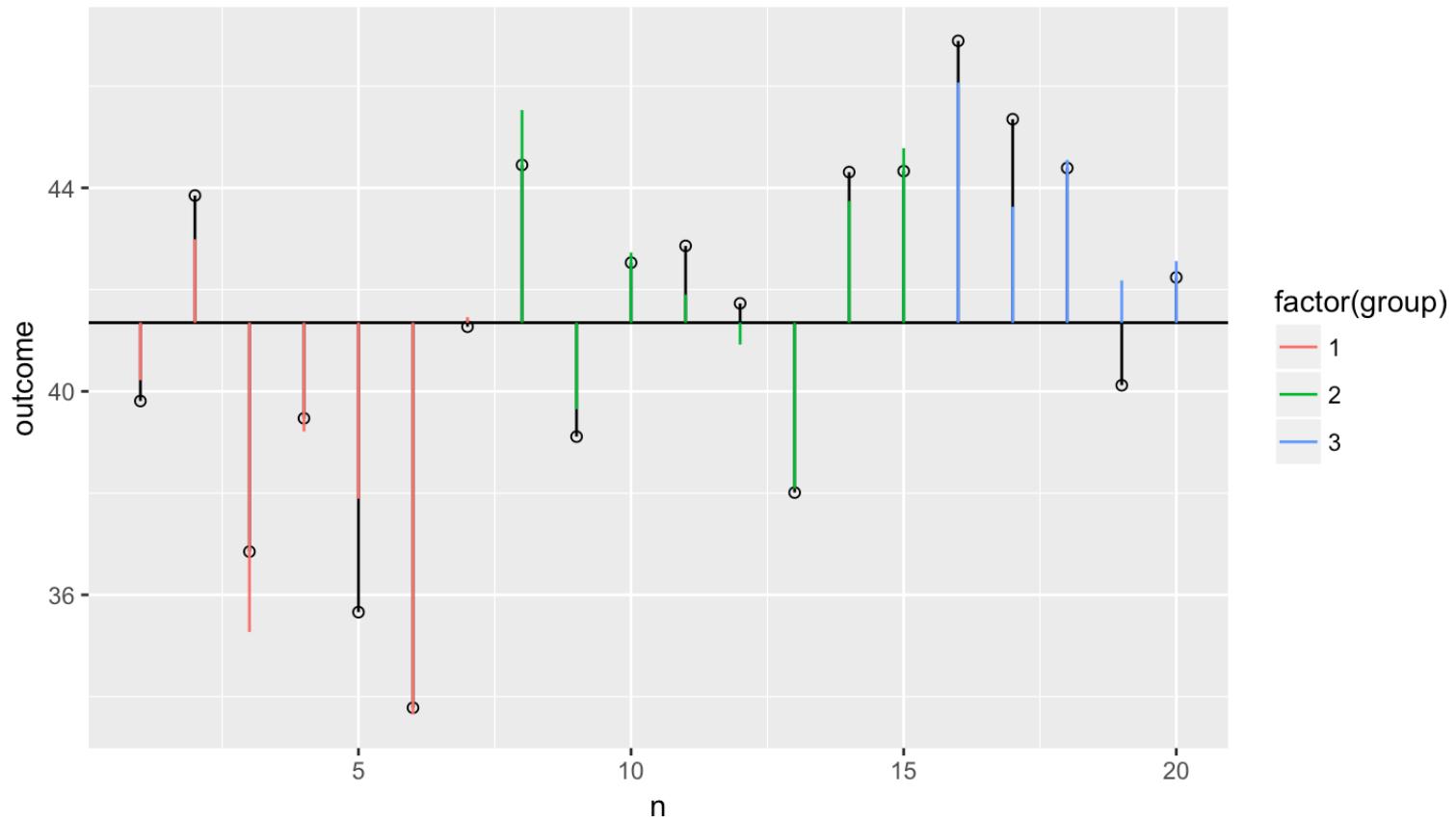


# Model variance group and covariate visual

```
plot + geom_segment(aes(x = n, y = grand.mean, xend = n, yend = data$model, co  
plot + ggtitle("Both group and covariate variance")
```



## Both group and covariate variance

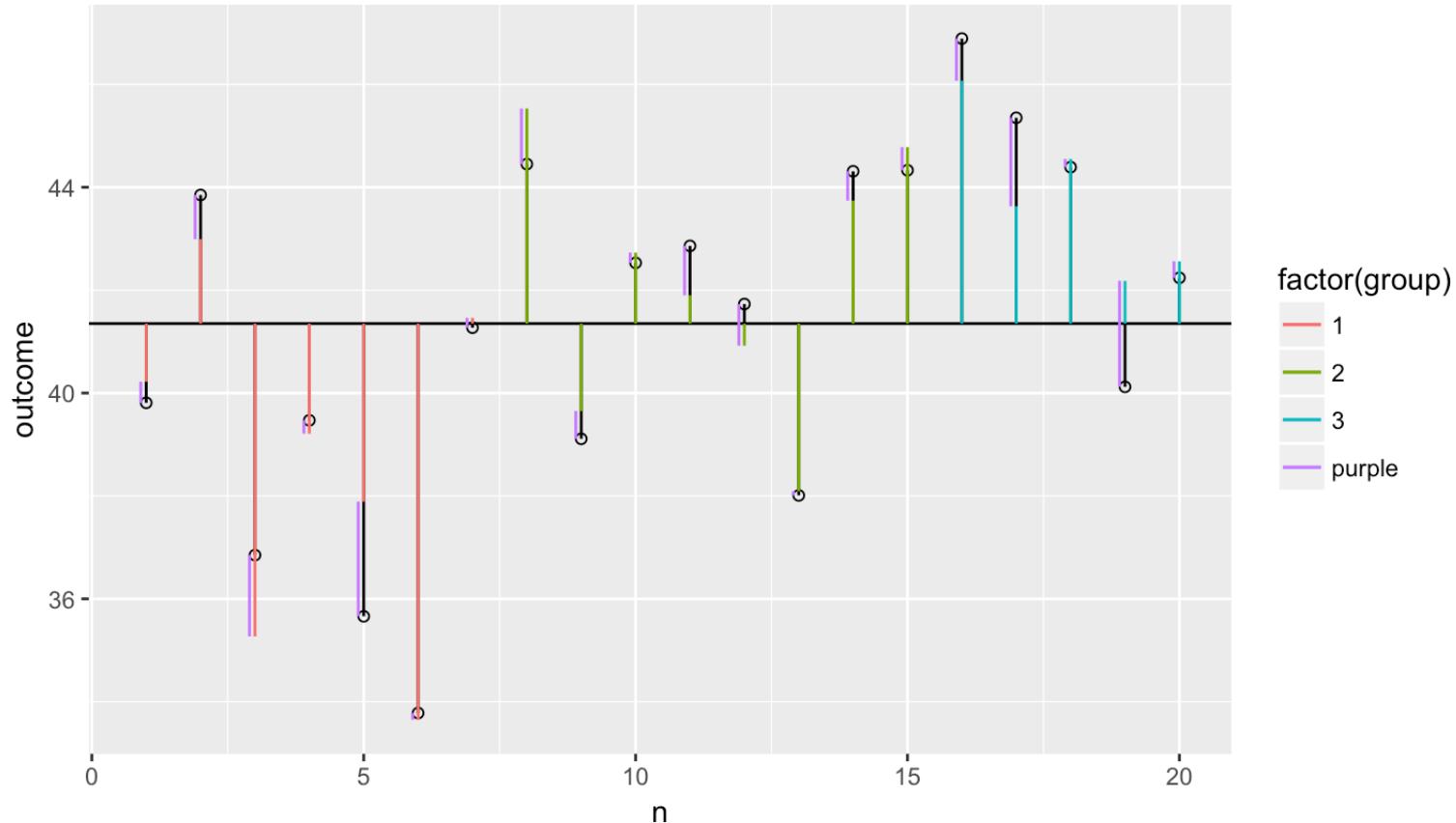


# Error variance with covariate

```
plot + geom_segment(aes(x      = n-.1,  
                         y      = outcome,  
                         xend   = n-.1,  
                         yend   = data$model,  
                         colour = "purple")) +  
  ggtitle("Error variance")
```



## Error variance



# Sums of squares

```
SS.model = with(data, sum((model - grand.mean)^2))
SS.error = with(data, sum((outcome - model)^2))

# Sums of squares for individual effects
SS.model.group = with(data, sum((model.group - grand.mean)^2))
SS.model.covar = with(data, sum((model.covar - grand.mean)^2))

SS.covar = SS.model - SS.model.group; SS.covar ## SS.covar corrected for group
```

```
## [1] 121.8463
```

```
SS.group = SS.model - SS.model.covar; SS.group ## SS.group corrected for covar
```

```
#  
[1] 54.65778
```

# F-ratio

$$F = \frac{MS_{model}}{MS_{error}} = \frac{SIGNAL}{NOISE}$$

```
n = 20
k = 3
df.model = k - 1
df.error = n - k - 1

MS.model = SS.group / df.model
MS.error = SS.error / df.error

F = MS.model / MS.error
F
```

```
## [1] 21.74406
```



# *P*-value

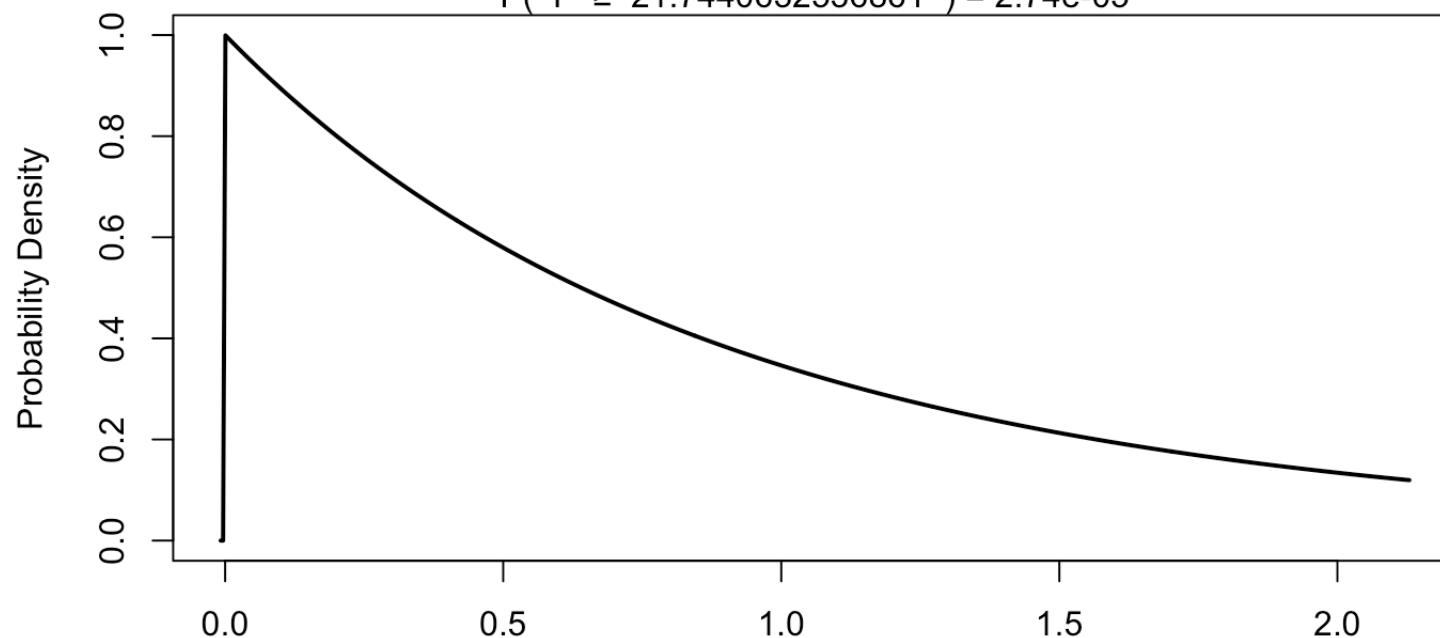
```
library("visualize")
visualize.f(F, df.model, df.error, section = "upper")
```



### F Distribution

**df1 = 2 df2 = 16**

$$P( F \geq 21.7440632536861 ) = 2.74e-05$$



F - Statistic

$$\mu = 1.14, \sigma^2 = 1.74$$

# Alpha & Power

```
F.values = seq(0, 30, .01)

plot(F.values, df(F.values, df.model, df.error), type = "l", ylab="density", m
critical.value = qf(.95, df.model, df.error)

critical.range = seq(critical.value, 30, .01)

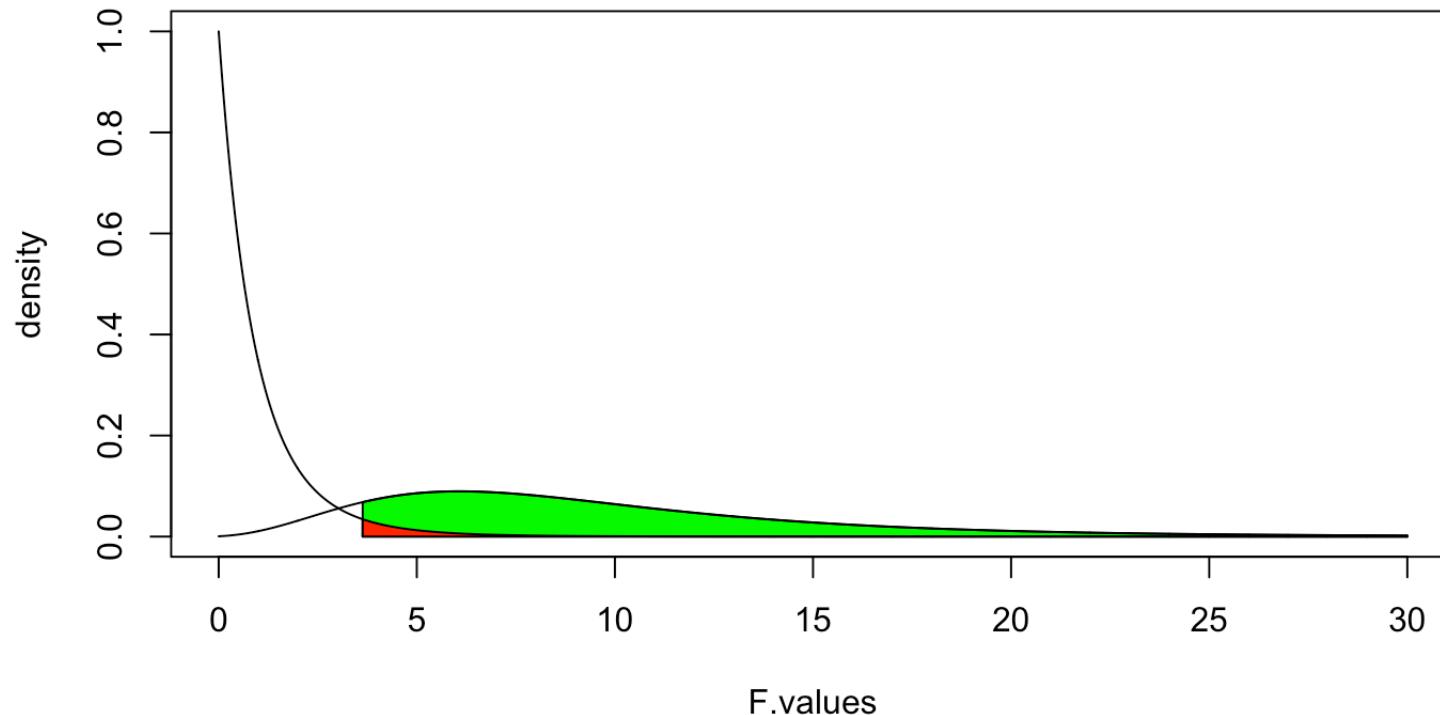
polygon(c(critical.range,rev(critical.range)),
         c(critical.range*0, rev(df(critical.range, df.model, df.error, ncp = 1

lines(F.values, df(F.values, df.model, df.error, ncp = 15))

polygon(c(critical.range,rev(critical.range)),
         c(critical.range*0, rev(df(critical.range, df.model, df.error))), col
```



## H0 and HA F-distribution



# Adjusted means

```
# Add dummy variables
data$dummy.1 <- ifelse(data$group == 2, 1, 0)
data$dummy.2 <- ifelse(data$group == 3, 1, 0)

# b coefficients
b.cov = fit$coefficients["covar"];           b.int = fit$coefficients["(Intercept)"]
b.2   = fit$coefficients["factor(group)2"]; b.3   = fit$coefficients["factor(group)3"]

# Adjustment factor for the means of the independent variable
data$mean.adj <- with(data, b.int + b.cov * mean(covar) + b.2 * dummy.1 + b.3 * dummy.2)

aggregate(mean.adj ~ group, data, mean)
```

```
##   group mean.adj
## 1      1 39.18576
## 2      2 41.95576
## 3      3 43.36576
```



# Real $\beta$ 's

```
b.0 = 15 # initial value for group 1  
b.1 = 3 # difference between group 1 and 2  
b.2 = 4 # difference between group 1 and 3  
b.3 = 3 # Weight for covariate  
  
cbind(m.covar = mu.covar*3,  
      BETA     = c(b.0, b.0+b.1, b.0+b.2),  
      sum       = mu.covar*3 + c(b.0, b.0+b.1, b.0+b.2))
```

```
##          m.covar BETA  sum  
## [1,]        24    15   39  
## [2,]        24    18   42  
## [3,]        24    19   43
```



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END