Data Analysis

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# **Information**

Please refer to ‘Data Cleaning’ script prior to accessing this script.

# **Setup**

knitr::opts\_chunk$set(echo = TRUE)  
require("knitr")

## Loading required package: knitr

opts\_knit$set(root.dir = "~/Library/Mobile Documents/com~apple~CloudDocs/Documents/Uni/Masters/Empirical Project/Code/Empirical\_Project")  
  
# turn off scientific notation  
options(scipen = 999)

### *Load Libraries*

library("ggplot2") # for figures  
library("psych") # for Cronbach's alpha, for describe function

##   
## Attaching package: 'psych'

## The following objects are masked from 'package:ggplot2':  
##   
## %+%, alpha

library("ppcor") # for partial correlation p-values

## Loading required package: MASS

library("dplyr") # for mutate function

##   
## Attaching package: 'dplyr'

## The following object is masked from 'package:MASS':  
##   
## select

## The following objects are masked from 'package:stats':  
##   
## filter, lag

## The following objects are masked from 'package:base':  
##   
## intersect, setdiff, setequal, union

library("ggpubr") # for qq-plots  
library("GGally") # for scatterplot matrix

## Registered S3 method overwritten by 'GGally':  
## method from   
## +.gg ggplot2

library("effsize") # for calculation of effect size

##   
## Attaching package: 'effsize'

## The following object is masked from 'package:psych':  
##   
## cohen.d

library("pwr") # for power calculation  
library("performance") # for assessing robustness of model  
library("effsize") # for eta squared  
library("reshape2") # for transforming data from wide to long format  
library("tidyverse") # for data cleaning

## ── Attaching packages ─────────────────────────────────────── tidyverse 1.3.1 ──

## ✓ tibble 3.1.3 ✓ purrr 0.3.4  
## ✓ tidyr 1.1.3 ✓ stringr 1.4.0  
## ✓ readr 2.0.0 ✓ forcats 0.5.1

## ── Conflicts ────────────────────────────────────────── tidyverse\_conflicts() ──  
## x psych::%+%() masks ggplot2::%+%()  
## x psych::alpha() masks ggplot2::alpha()  
## x dplyr::filter() masks stats::filter()  
## x dplyr::lag() masks stats::lag()  
## x dplyr::select() masks MASS::select()

library("rstatix") # for ANOVA and ANCOVA

##   
## Attaching package: 'rstatix'

## The following object is masked from 'package:MASS':  
##   
## select

## The following object is masked from 'package:stats':  
##   
## filter

library("gridExtra") # for grid.arrange function

##   
## Attaching package: 'gridExtra'

## The following object is masked from 'package:dplyr':  
##   
## combine

library("car") # for levene's test

## Loading required package: carData

##   
## Attaching package: 'car'

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

## The following object is masked from 'package:dplyr':  
##   
## recode

## The following object is masked from 'package:psych':  
##   
## logit

library("emmeans") # to obtain estimated marginal means

##   
## Attaching package: 'emmeans'

## The following object is masked from 'package:GGally':  
##   
## pigs

### *Set Working Directory*

# please change this to your own working directory path  
setwd("~/Library/Mobile Documents/com~apple~CloudDocs/Documents/Uni/Masters/Empirical Project/Code/Empirical\_Project")

### *Read in Data and Save Data to an Object*

# please change this to however you have stored the data file   
# reading in dataframe 2, as this is the one with exclusion of n = 5  
df <- read.csv(file = "data/cleaned/dataframe\_2.csv", header = TRUE, na.strings = "NA")

### *Change Variable Classifications*

# change variable classifications to meet requirements for later analyses  
# ensure IVs and categorical variables are factor variables  
# and DVs or continuous variables are numeric variables  
  
# participant id and demographics  
df$id <- factor(df$id)  
df$age <- as.numeric(df$age)  
df$sex <- factor(df$sex)  
df$ethnicity <- factor(df$ethnicity)  
df$sexual\_orientation <- factor(df$sexual\_orientation)  
  
# fixation count DVs  
df$acq\_csp\_fix\_count <- as.numeric(df$acq\_csp\_fix\_count)  
df$acq\_csm\_fix\_count <- as.numeric(df$acq\_csm\_fix\_count)  
df$ext\_csp\_fix\_count <- as.numeric(df$ext\_csp\_fix\_count)  
df$ext\_csm\_fix\_count <- as.numeric(df$ext\_csm\_fix\_count)  
df$e\_ext\_csp\_fix\_count <- as.numeric(df$e\_ext\_csp\_fix\_count)  
df$l\_ext\_csp\_fix\_count <- as.numeric(df$l\_ext\_csp\_fix\_count)  
df$e\_ext\_csm\_fix\_count <- as.numeric(df$e\_ext\_csm\_fix\_count)  
df$l\_ext\_csm\_fix\_count <- as.numeric(df$l\_ext\_csm\_fix\_count)  
  
# fixation duration DVs  
df$acq\_csp\_fix\_duration <- as.numeric(df$acq\_csp\_fix\_duration)  
df$acq\_csm\_fix\_duration <- as.numeric(df$acq\_csm\_fix\_duration)  
df$ext\_csp\_fix\_duration <- as.numeric(df$ext\_csp\_fix\_duration)  
df$ext\_csm\_fix\_duration <- as.numeric(df$ext\_csm\_fix\_duration)  
df$e\_ext\_csp\_fix\_duration <- as.numeric(df$e\_ext\_csp\_fix\_duration)  
df$l\_ext\_csp\_fix\_duration <- as.numeric(df$l\_ext\_csp\_fix\_duration)  
df$e\_ext\_csm\_fix\_duration <- as.numeric(df$e\_ext\_csm\_fix\_duration)  
df$l\_ext\_csm\_fix\_duration <- as.numeric(df$l\_ext\_csm\_fix\_duration)  
  
# saccade amplitude DVs  
df$acq\_csp\_sacc\_amplitude <- as.numeric(df$acq\_csp\_sacc\_amplitude)  
df$acq\_csm\_sacc\_amplitude <- as.numeric(df$acq\_csm\_sacc\_amplitude)  
df$ext\_csp\_sacc\_amplitude <- as.numeric(df$ext\_csp\_sacc\_amplitude)  
df$ext\_csm\_sacc\_amplitude <- as.numeric(df$ext\_csm\_sacc\_amplitude)  
df$e\_ext\_csp\_sacc\_amplitude <- as.numeric(df$e\_ext\_csp\_sacc\_amplitude)  
df$l\_ext\_csp\_sacc\_amplitude <- as.numeric(df$l\_ext\_csp\_sacc\_amplitude)  
df$e\_ext\_csm\_sacc\_amplitude <- as.numeric(df$e\_ext\_csm\_sacc\_amplitude)  
df$l\_ext\_csm\_sacc\_amplitude <- as.numeric(df$l\_ext\_csm\_sacc\_amplitude)

# **Internal Consistency of IUS and STICSA**

## IUS total  
# compute & extract alpha value and save as an object  
alpha\_ius <- psych::alpha(df[, c("ius\_1", "ius\_2", "ius\_3", "ius\_4",  
 "ius\_5", "ius\_6", "ius\_7", "ius\_8",  
 "ius\_9", "ius\_10", "ius\_11", "ius\_12",  
 "ius\_13", "ius\_14", "ius\_15", "ius\_16",  
 "ius\_17", "ius\_18", "ius\_19", "ius\_20",   
 "ius\_21", "ius\_22", "ius\_23", "ius\_24",  
 "ius\_25", "ius\_26", "ius\_27")])$total[1]  
  
## STICSA total  
# compute & extract alpha value and save as an object  
alpha\_sticsa <- psych::alpha(df[, c("sticsa\_1", "sticsa\_2", "sticsa\_3", "sticsa\_4",  
 "sticsa\_5", "sticsa\_6", "sticsa\_7", "sticsa\_8",  
 "sticsa\_9", "sticsa\_10", "sticsa\_11", "sticsa\_12",  
 "sticsa\_13", "sticsa\_14", "sticsa\_15", "sticsa\_16",  
 "sticsa\_17", "sticsa\_18", "sticsa\_19", "sticsa\_20",   
 "sticsa\_21")])$total[1]  
  
# create table of both Crobach's alpha values  
cronbachs\_alpha\_questionnaires <- rbind(alpha\_ius, alpha\_sticsa)  
  
# clean up row and column names for easier interpretation  
rownames(cronbachs\_alpha\_questionnaires) <- c("IUS-27", "STICSA")  
colnames(cronbachs\_alpha\_questionnaires) <- "Cronbach's Alpha"  
  
# obtain Cronbach's alpha table  
cronbachs\_alpha\_questionnaires

## Cronbach's Alpha  
## IUS-27 0.9496736  
## STICSA 0.8766597

# *Compute Questionnaire Totals*

#### IUS total  
# all items, no reverse scoring  
df$ius\_total <- as.numeric(df$ius\_1 + df$ius\_2 + df$ius\_3 + df$ius\_4 + df$ius\_5 +   
 df$ius\_6 + df$ius\_7 + df$ius\_8 + df$ius\_9 +   
 df$ius\_10 + df$ius\_11 + df$ius\_12 + df$ius\_13 +   
 df$ius\_14 + df$ius\_15 + df$ius\_16 + df$ius\_17 +   
 df$ius\_18 + df$ius\_19 + df$ius\_20 + df$ius\_21 +   
 df$ius\_22 + df$ius\_23 + df$ius\_24 + df$ius\_25 +   
 df$ius\_26 + df$ius\_27)  
  
#### STICSA total  
# all items, no reverse scoring  
df$sticsa\_total <- as.numeric(df$sticsa\_1 + df$sticsa\_2 + df$sticsa\_3 +  
 df$sticsa\_4 + df$sticsa\_5 + df$sticsa\_6 +  
 df$sticsa\_7 + df$sticsa\_8 + df$sticsa\_9 +  
 df$sticsa\_10 + df$sticsa\_11 + df$sticsa\_12 +  
 df$sticsa\_13 + df$sticsa\_14 + df$sticsa\_15 +  
 df$sticsa\_16 + df$sticsa\_17 + df$sticsa\_18 +  
 df$sticsa\_19 + df$sticsa\_20 + df$sticsa\_21)

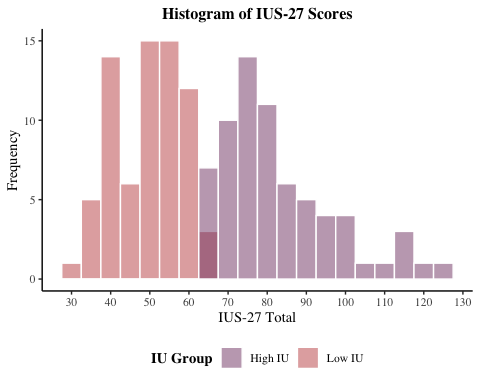
# *Create High / Low IU Classifications*

# compute variable classifying participants as high/ low IU on basis of median split,   
# and store as factor  
df$iu\_group <- factor(ifelse(df$ius\_total >= 65, 1, -1))  
# high IU = 1  
# low IU = -1

# *Check Distribution and Range to Identify Extreme Scores and Potential Data Errors in Questionnaires*

## *For IUS 27 Total in Both Groups*

# possible total scores for the IUS range from 27-135   
  
########################## check distributions   
hist\_ius\_total <- df %>%  
 ggplot(aes(ius\_total, fill = iu\_group)) +  
 geom\_histogram(binwidth = 5, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(20, 140, 10)) +  
 labs(x = "IUS-27 Total", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram of IUS-27 Scores") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_ius\_total



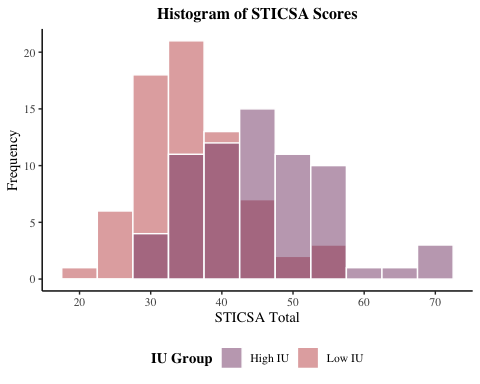
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_ius\_total.png",   
 plot = hist\_ius\_total,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
########################## check ranges  
range\_ius\_total <- by(df$ius\_total, df$iu\_group, range)  
range\_ius\_total

## df$iu\_group: -1  
## [1] 32 64  
## ------------------------------------------------------------   
## df$iu\_group: 1  
## [1] 65 125

# for high IU: 65-125  
# for low IU: 32-64  
##### overall: all scores are in range of possible scores, no errors apparent

## *For STICSA Total in Both Groups*

# possible total scores for the STICSA range from 21-84  
  
########################## check distributions   
hist\_sticsa\_total <- df %>%  
 ggplot(aes(sticsa\_total, fill = iu\_group)) +  
 geom\_histogram(binwidth = 5, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(20, 90, 10)) +   
 labs(x = "STICSA Total", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram of STICSA Scores") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_sticsa\_total



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_sticsa\_total.png",   
 plot = hist\_sticsa\_total,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
########################## check ranges  
range\_sticsa\_total <- by(df$sticsa\_total, df$iu\_group, range)  
range\_sticsa\_total

## df$iu\_group: -1  
## [1] 22 57  
## ------------------------------------------------------------   
## df$iu\_group: 1  
## [1] 30 69

# for high IU: 30-69  
# for low IU: 22-57  
##### overall: all scores are in range of possible scores, no errors apparent

# **Compute Demographics**

#### for age  
# for all participants  
all\_age\_table <-   
 describe(df[, "age"])  
  
# for high IU  
high\_iu\_age\_table <-   
 describe(df[df$iu\_group =="1", "age"])  
  
# for low IU  
low\_iu\_age\_table <-   
 describe(df[df$iu\_group =="-1", "age"])  
  
# combine in a table  
age\_table <- rbind(all\_age\_table, high\_iu\_age\_table, low\_iu\_age\_table)  
  
# re-name rows for easier interpretation  
rownames(age\_table) <- c("Age (All Participants","Age (High IU Group)",   
 "Age (Low IU Group)")  
  
### for sex   
sex\_table <- xtabs(~ iu\_group + sex, data = df)  
sex\_table <- prop.table(sex\_table) %>%  
 round(digits = 4) \* 100  
rownames(sex\_table) <- c("Low IU", "High IU")  
sex\_table

## sex  
## iu\_group Female Male  
## Low IU 26.28 24.82  
## High IU 34.31 14.60

### for sexual orientation  
sexual\_orientation\_table <- xtabs(~ iu\_group + sexual\_orientation, data = df)  
sexual\_orientation\_table <- prop.table(sexual\_orientation\_table) %>%  
 round(digits = 4) \* 100  
rownames(sexual\_orientation\_table) <- c("Low IU", "High IU")  
sexual\_orientation\_table

## sexual\_orientation  
## iu\_group Heterosexual Sexual Minority  
## Low IU 42.15 7.44  
## High IU 42.98 7.44

### for ethnicity  
ethnicity\_table <- xtabs(~ iu\_group + ethnicity, data = df)  
ethnicity\_table <- prop.table(ethnicity\_table) %>%  
 round(digits = 4) \* 100  
rownames(ethnicity\_table) <- c("Low IU", "High IU")  
ethnicity\_table

## ethnicity  
## iu\_group Asian Black Middle Eastern/ Arab Mixed White  
## Low IU 7.26 1.61 2.42 0.81 37.90  
## High IU 16.13 0.00 0.81 0.81 32.26

#### write each to csv   
# age  
write.csv(age\_table, file = "tables/demographics/age\_table.csv",  
 row.names = TRUE)  
  
# ethnicity  
write.csv(ethnicity\_table, file = "tables/demographics/ethnicity\_table.csv",  
 row.names = TRUE)  
  
# sex  
write.csv(sex\_table, file = "tables/demographics/sex\_table.csv",   
 row.names = TRUE)  
  
# sexual orientation  
write.csv(sexual\_orientation\_table, file = "tables/demographics/sexual\_orientation\_table.csv",  
 row.names = TRUE)

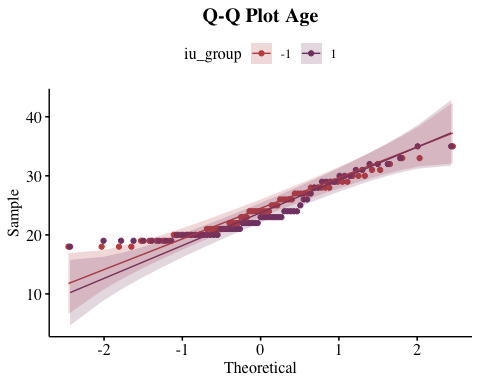
# *Check for Difference in Demographics Between Groups*

## *Check for Difference in Age Between Groups*

# t-test to check for intergroup differences in age  
  
# first check assumptions of t-test  
# plot data for both groups using QQ plot  
qqplot\_age <- ggqqplot(df, x = "age",  
 color = "iu\_group",  
 palette = c("#c45150", "#824372"),  
 title = "Q-Q Plot Age") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15))   
# inspect the QQ plots  
qqplot\_age

## Warning: Removed 1 rows containing non-finite values (stat\_qq).

## Warning: Removed 1 rows containing non-finite values (stat\_qq\_line).  
  
## Warning: Removed 1 rows containing non-finite values (stat\_qq\_line).



# save plot to file  
ggsave(filename = "graphs/qqplots/qqplot\_age.png",   
 plot = qqplot\_age,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

## Warning: Removed 1 rows containing non-finite values (stat\_qq).  
  
## Warning: Removed 1 rows containing non-finite values (stat\_qq\_line).  
  
## Warning: Removed 1 rows containing non-finite values (stat\_qq\_line).

# check significance of data for both groups using Shapiro-Wilk Test  
shapiro\_age <- by(df$age, df$iu\_group, shapiro.test)  
shapiro\_age

## df$iu\_group: -1  
##   
## Shapiro-Wilk normality test  
##   
## data: dd[x, ]  
## W = 0.95698, p-value = 0.016  
##   
## ------------------------------------------------------------   
## df$iu\_group: 1  
##   
## Shapiro-Wilk normality test  
##   
## data: dd[x, ]  
## W = 0.88408, p-value = 0.00001422

# high IU: p-value < .05, data violate assumption of normality  
# low IU: p-value < .05, data violate assumption of normality  
  
## check assumption of homogeneity of variances using Bartlett Test ##  
bartlett\_age <- bartlett.test(age ~ iu\_group, data = df)   
bartlett\_age

##   
## Bartlett test of homogeneity of variances  
##   
## data: age by iu\_group  
## Bartlett's K-squared = 0.27665, df = 1, p-value = 0.5989

# p-value > .05, data meet assumption of equal variances  
  
## compute independent samples t.test ##  
# as data violate assumption of normality,  
# use non-parametric Mann Whitney U  
  
# compute t.test and assign values to an object  
age\_groupdiff <- wilcox.test(age ~ iu\_group, data = df, paired = FALSE)  
  
# obtain t.test values   
age\_groupdiff

##   
## Wilcoxon rank sum test with continuity correction  
##   
## data: age by iu\_group  
## W = 2585.5, p-value = 0.3773  
## alternative hypothesis: true location shift is not equal to 0

# p-value > .05, there is no statistical difference in age between groups

## *Check for Difference in Ethnicity Between Groups*

# compute chi-square of cross-tabulation and save as object   
chi\_ethnicity <- chisq.test(table(df$iu\_group, df$ethnicity))

## Warning in chisq.test(table(df$iu\_group, df$ethnicity)): Chi-squared  
## approximation may be incorrect

# check assumption of chi-square  
chi\_ethnicity$expected

##   
## Asian Black Middle Eastern/ Arab Mixed White  
## -1 14.5 1 2 1 43.5  
## 1 14.5 1 2 1 43.5

# multiple cells with values less than 5, does not meet assumptions  
# and therefore requires Fisher's Exact Test  
  
# obtain statistic and df  
chi\_ethnicity

##   
## Pearson's Chi-squared test  
##   
## data: table(df$iu\_group, df$ethnicity)  
## X-squared = 7.7356, df = 4, p-value = 0.1018

# obtain corrected p-value   
chi\_ethnicity\_pval <- fisher.test(df$iu\_group, df$ethnicity)   
chi\_ethnicity\_pval

##   
## Fisher's Exact Test for Count Data  
##   
## data: df$iu\_group and df$ethnicity  
## p-value = 0.05899  
## alternative hypothesis: two.sided

# p-value > .05, no evidence of statistical difference in ethnicity between groups

## *Check for Difference in Sex Between Groups*

# compute chi-square of cross-tabulation and save as object   
chi\_sex <- chisq.test(table(df$iu\_group, df$sex))  
  
# check assumption of chi-square  
chi\_sex\_expected <- chi\_sex$expected   
chi\_sex\_expected

##   
## Female Male  
## -1 42.40876 27.59124  
## 1 40.59124 26.40876

# no cells less than 5, meets assumptions   
  
# obtain statistic, df and p-value  
chi\_sex

##   
## Pearson's Chi-squared test with Yates' continuity correction  
##   
## data: table(df$iu\_group, df$sex)  
## X-squared = 4.2708, df = 1, p-value = 0.03877

# p-value < .05, there appears to be a statistical difference in sex between groups  
  
# therefore, obtain observed values  
chi\_sex\_observed <- chi\_sex$observed  
chi\_sex\_observed

##   
## Female Male  
## -1 36 34  
## 1 47 20

## *Check for Difference in Sexual Orientation Between Groups*

# compute chi-square of cross-tabulation and save as object   
chi\_sexual\_orientation <- chisq.test(table(df$iu\_group, df$sexual\_orientation))   
  
# check assumption of chi-square  
chi\_sexual\_orientation$expected

##   
## Heterosexual Sexual Minority  
## -1 51.07438 8.92562  
## 1 51.92562 9.07438

# no cells with values less than 5, meets assumptions  
  
# obtain statistic and df  
chi\_sexual\_orientation

##   
## Pearson's Chi-squared test with Yates' continuity correction  
##   
## data: table(df$iu\_group, df$sexual\_orientation)  
## X-squared = 0, df = 1, p-value = 1

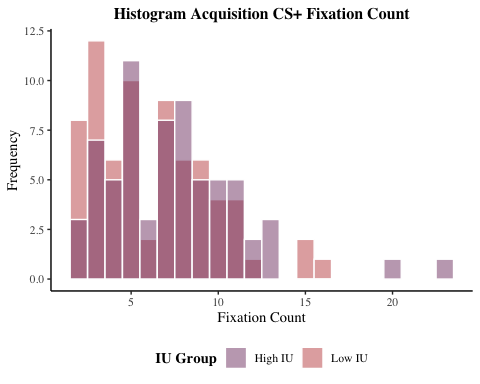
# p-value > .05, no evidence of statistical difference in sexual orientation between groups

# *Distribution Checks of Eye-Movement Variables*

## **Fixation Count**

### *Acquisition CS+*

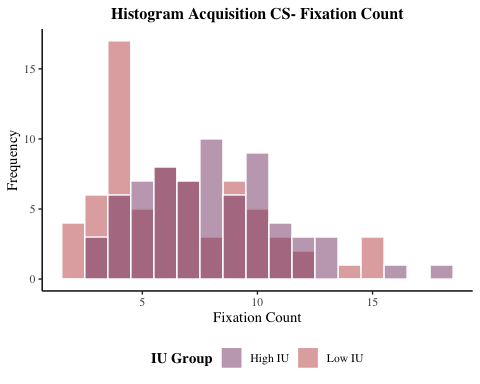
hist\_acq\_csp\_fix\_count <- df %>%  
 ggplot(aes(acq\_csp\_fix\_count, fill = iu\_group)) +  
 geom\_histogram(binwidth = 1, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 30, 5)) +  
 labs(x = "Fixation Count", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Acquisition CS+ Fixation Count") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_acq\_csp\_fix\_count



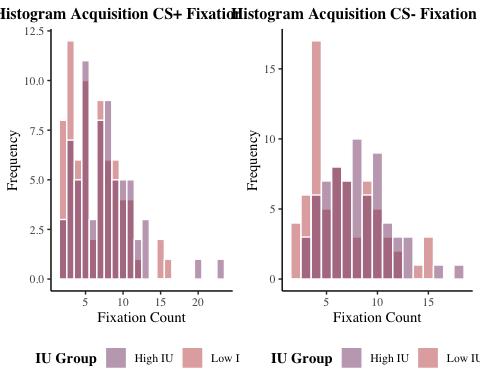
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_acq\_csp\_fix\_count.png",   
 plot = hist\_acq\_csp\_fix\_count,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Acquisition CS-*

hist\_acq\_csm\_fix\_count <- df %>%  
 ggplot(aes(acq\_csm\_fix\_count, fill = iu\_group)) +  
 geom\_histogram(binwidth = 1, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 30, 5)) +  
 labs(x = "Fixation Count", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Acquisition CS- Fixation Count") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_acq\_csm\_fix\_count



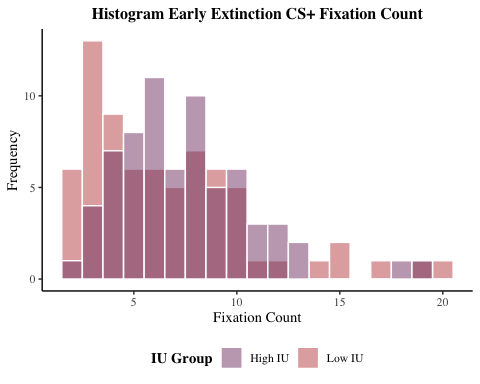
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_acq\_csm\_fix\_count.png",   
 plot = hist\_acq\_csm\_fix\_count,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
# combine acquisition fixation count graphs  
hists\_acq\_fix\_count <-   
 grid.arrange(hist\_acq\_csp\_fix\_count, hist\_acq\_csm\_fix\_count,  
 ncol =2)



# save plot to file  
ggsave(filename = "graphs/histograms/hists\_acq\_fix\_count.png",   
 plot = hists\_acq\_fix\_count,   
 width = 30,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Early Extinction CS+*

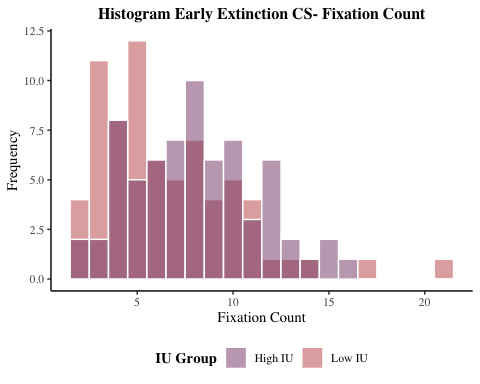
hist\_e\_ext\_csp\_fix\_count <- df %>%  
 ggplot(aes(e\_ext\_csp\_fix\_count, fill = iu\_group)) +  
 geom\_histogram(binwidth = 1, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 30, 5)) +  
 labs(x = "Fixation Count", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Early Extinction CS+ Fixation Count") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_e\_ext\_csp\_fix\_count



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_e\_ext\_csp\_fix\_count.png",   
 plot = hist\_e\_ext\_csp\_fix\_count,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Early Extinction CS-*

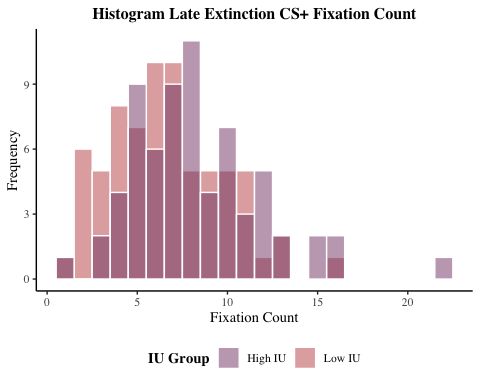
hist\_e\_ext\_csm\_fix\_count <- df %>%  
 ggplot(aes(e\_ext\_csm\_fix\_count, fill = iu\_group)) +  
 geom\_histogram(binwidth = 1, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 30, 5)) +  
 labs(x = "Fixation Count", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Early Extinction CS- Fixation Count") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_e\_ext\_csm\_fix\_count



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_e\_ext\_csm\_fix\_count.png",   
 plot = hist\_e\_ext\_csm\_fix\_count,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Late Extinction CS+*

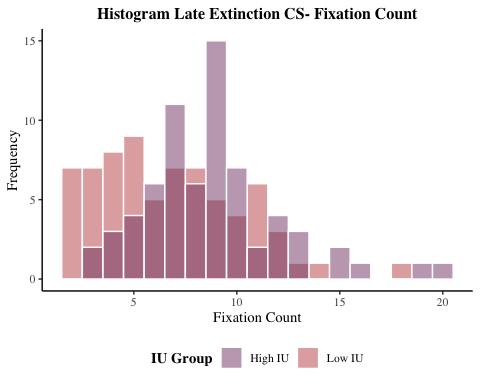
hist\_l\_ext\_csp\_fix\_count <- df %>%  
 ggplot(aes(l\_ext\_csp\_fix\_count, fill = iu\_group)) +  
 geom\_histogram(binwidth = 1, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 30, 5)) +  
 labs(x = "Fixation Count", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Late Extinction CS+ Fixation Count") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_l\_ext\_csp\_fix\_count



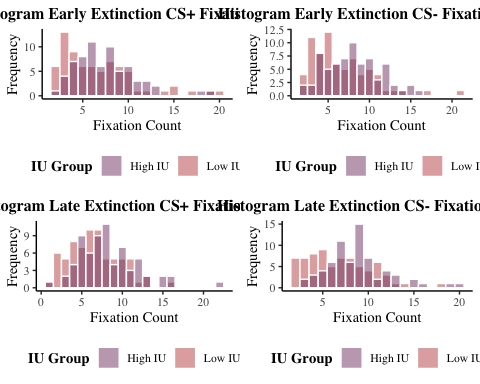
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_l\_ext\_csp\_fix\_count.png",   
 plot = hist\_l\_ext\_csp\_fix\_count,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Late Extinction CS-*

hist\_l\_ext\_csm\_fix\_count <- df %>%  
 ggplot(aes(l\_ext\_csm\_fix\_count, fill = iu\_group)) +  
 geom\_histogram(binwidth = 1, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 30, 5)) +  
 labs(x = "Fixation Count", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Late Extinction CS- Fixation Count") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_l\_ext\_csm\_fix\_count



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_l\_ext\_csm\_fix\_count.png",   
 plot = hist\_l\_ext\_csm\_fix\_count,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
# combine extinction fixation count graphs  
hists\_ext\_fix\_count <-   
 grid.arrange(hist\_e\_ext\_csp\_fix\_count, hist\_e\_ext\_csm\_fix\_count,  
 hist\_l\_ext\_csp\_fix\_count, hist\_l\_ext\_csm\_fix\_count,  
 ncol =2)

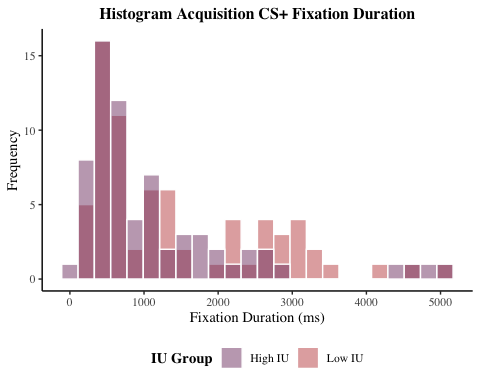


# save plot to file  
ggsave(filename = "graphs/histograms/hists\_ext\_fix\_count.png",   
 plot = hists\_ext\_fix\_count,   
 width = 30,   
 height = 20,   
 dpi = 300,   
 units = "cm")

## **Fixation Duration**

### *Acquisition CS+*

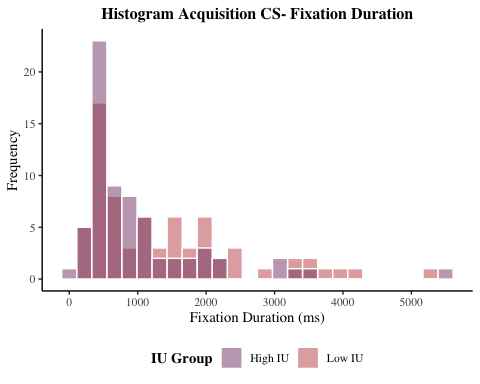
hist\_acq\_csp\_fix\_duration <- df %>%  
 ggplot(aes(acq\_csp\_fix\_duration, fill = iu\_group)) +  
 geom\_histogram(binwidth = 220, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 6000, 1000)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Acquisition CS+ Fixation Duration") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_acq\_csp\_fix\_duration



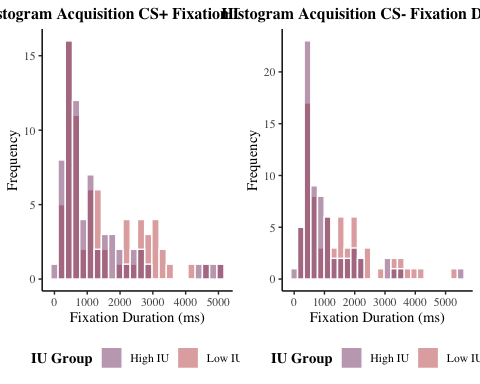
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_acq\_csp\_fix\_duration.png",   
 plot = hist\_acq\_csp\_fix\_duration,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Acquisition CS-*

hist\_acq\_csm\_fix\_duration <- df %>%  
 ggplot(aes(acq\_csm\_fix\_duration, fill = iu\_group)) +  
 geom\_histogram(binwidth = 220, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 6000, 1000)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Acquisition CS- Fixation Duration") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_acq\_csm\_fix\_duration



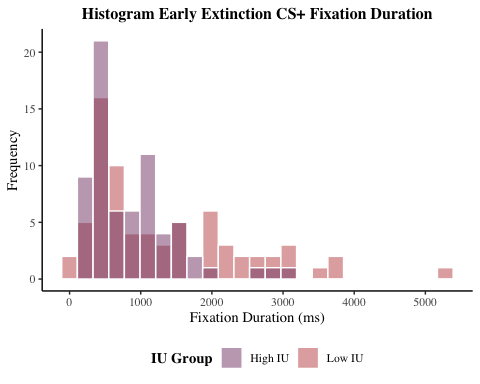
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_acq\_csm\_fix\_duration.png",   
 plot = hist\_acq\_csm\_fix\_duration,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
# combine acquisition fixation duration graphs  
hists\_acq\_fix\_duration <-   
 grid.arrange(hist\_acq\_csp\_fix\_duration, hist\_acq\_csm\_fix\_duration,  
 ncol =2)



# save plot to file  
ggsave(filename = "graphs/histograms/hists\_acq\_fix\_duration.png",   
 plot = hists\_acq\_fix\_duration,   
 width = 30,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Early Extinction CS+*

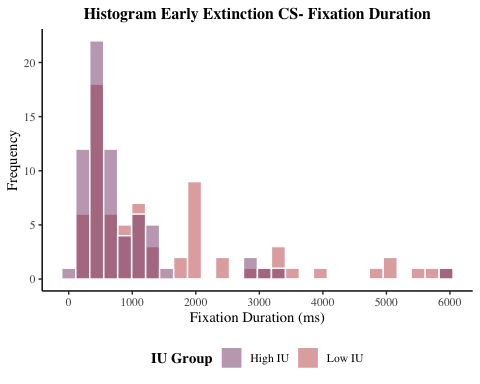
hist\_e\_ext\_csp\_fix\_duration <- df %>%  
 ggplot(aes(e\_ext\_csp\_fix\_duration, fill = iu\_group)) +  
 geom\_histogram(binwidth = 220, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 6000, 1000)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Early Extinction CS+ Fixation Duration") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_e\_ext\_csp\_fix\_duration



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_e\_ext\_csp\_fix\_duration.png",   
 plot = hist\_e\_ext\_csp\_fix\_duration,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Early Extinction CS-*

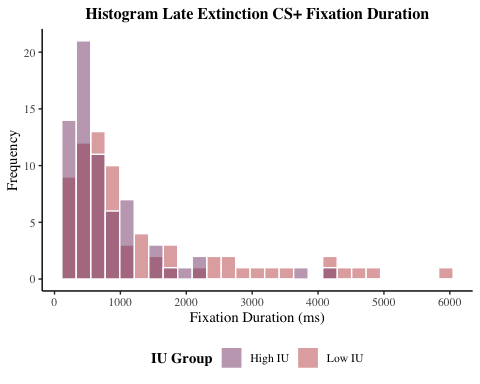
hist\_e\_ext\_csm\_fix\_duration <- df %>%  
 ggplot(aes(e\_ext\_csm\_fix\_duration, fill = iu\_group)) +  
 geom\_histogram(binwidth = 220, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 6000, 1000)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Early Extinction CS- Fixation Duration") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_e\_ext\_csm\_fix\_duration



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_e\_ext\_csm\_fix\_duration.png",   
 plot = hist\_e\_ext\_csm\_fix\_duration,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Late Extinction CS+*

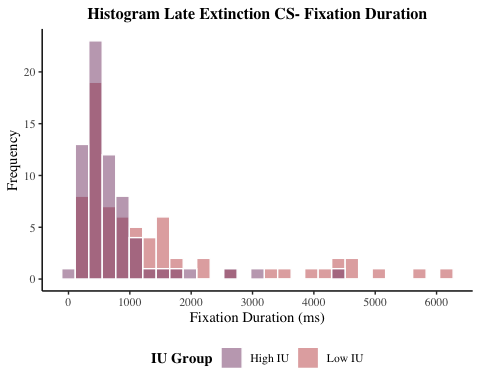
hist\_l\_ext\_csp\_fix\_duration <- df %>%  
 ggplot(aes(l\_ext\_csp\_fix\_duration, fill = iu\_group)) +  
 geom\_histogram(binwidth = 220, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 6000, 1000)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Late Extinction CS+ Fixation Duration") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_l\_ext\_csp\_fix\_duration



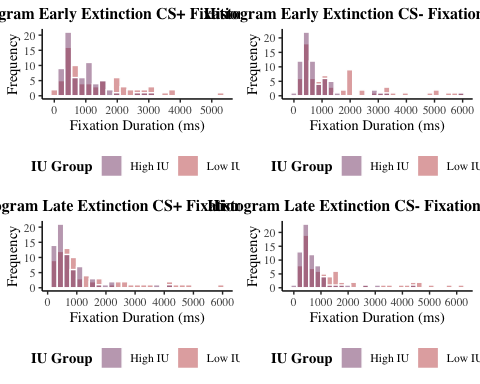
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_l\_ext\_csp\_fix\_duration.png",   
 plot = hist\_l\_ext\_csp\_fix\_duration,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Late Extinction CS-*

hist\_l\_ext\_csm\_fix\_duration <- df %>%  
 ggplot(aes(l\_ext\_csm\_fix\_duration, fill = iu\_group)) +  
 geom\_histogram(binwidth = 220, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 6000, 1000)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Late Extinction CS- Fixation Duration") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_l\_ext\_csm\_fix\_duration



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_l\_ext\_csm\_fix\_duration.png",   
 plot = hist\_l\_ext\_csm\_fix\_duration,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
# combine extinction fixation duration graphs  
hists\_ext\_fix\_duration <-   
 grid.arrange(hist\_e\_ext\_csp\_fix\_duration, hist\_e\_ext\_csm\_fix\_duration,  
 hist\_l\_ext\_csp\_fix\_duration, hist\_l\_ext\_csm\_fix\_duration,  
 ncol =2)

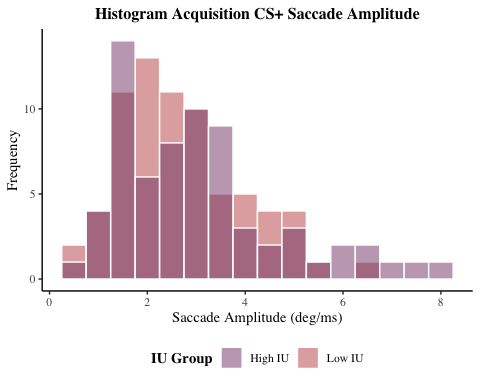


# save plot to file  
ggsave(filename = "graphs/histograms/hists\_ext\_fix\_duration.png",   
 plot = hists\_ext\_fix\_duration,   
 width = 30,   
 height = 20,   
 dpi = 300,   
 units = "cm")

## **Saccade Amplitude**

### *Acquisition CS+*

hist\_acq\_csp\_sacc\_amplitude <- df %>%  
 ggplot(aes(acq\_csp\_sacc\_amplitude, fill = iu\_group)) +  
 geom\_histogram(binwidth = .5, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 10, 2)) +  
 labs(x = "Saccade Amplitude (deg/ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Acquisition CS+ Saccade Amplitude") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_acq\_csp\_sacc\_amplitude

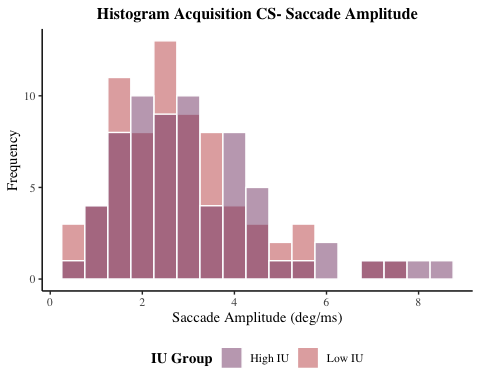


ggsave(filename = "graphs/histograms/hist\_acq\_csp\_sacc\_amplitude.png",   
 plot = hist\_acq\_csp\_sacc\_amplitude,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Acquisition CS-*

hist\_acq\_csm\_sacc\_amplitude <- df %>%  
 ggplot(aes(acq\_csm\_sacc\_amplitude, fill = iu\_group)) +  
 geom\_histogram(binwidth = .5, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 10, 2)) +  
 labs(x = "Saccade Amplitude (deg/ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Acquisition CS- Saccade Amplitude") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_acq\_csm\_sacc\_amplitude

## Warning: Removed 2 rows containing non-finite values (stat\_bin).

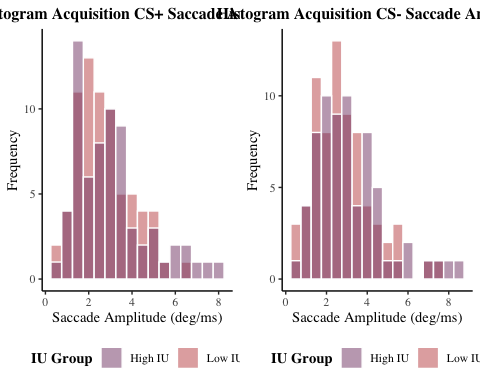


ggsave(filename = "graphs/histograms/hist\_acq\_csm\_sacc\_amplitude.png",   
 plot = hist\_acq\_csm\_sacc\_amplitude,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

## Warning: Removed 2 rows containing non-finite values (stat\_bin).

# combine acquisition saccade amplitude graphs  
hists\_acq\_sacc\_amplitude <-   
 grid.arrange(hist\_acq\_csp\_sacc\_amplitude, hist\_acq\_csm\_sacc\_amplitude,  
 ncol =2)

## Warning: Removed 2 rows containing non-finite values (stat\_bin).

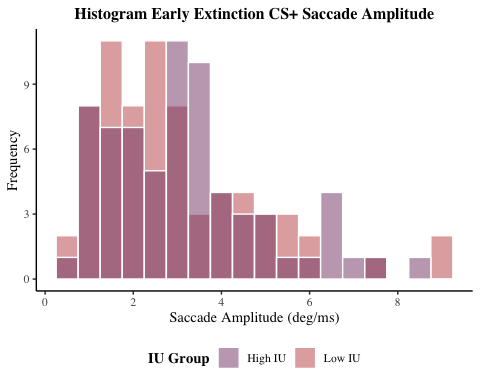


# save plot to file  
ggsave(filename = "graphs/histograms/hists\_acq\_sacc\_amplitude.png",   
 plot = hists\_acq\_sacc\_amplitude,   
 width = 30,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Early Extinction CS+*

hist\_e\_ext\_csp\_sacc\_amplitude <- df %>%  
 ggplot(aes(e\_ext\_csp\_sacc\_amplitude, fill = iu\_group)) +  
 geom\_histogram(binwidth = .5, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 14, 2)) +  
 labs(x = "Saccade Amplitude (deg/ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Early Extinction CS+ Saccade Amplitude") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_e\_ext\_csp\_sacc\_amplitude

## Warning: Removed 1 rows containing non-finite values (stat\_bin).



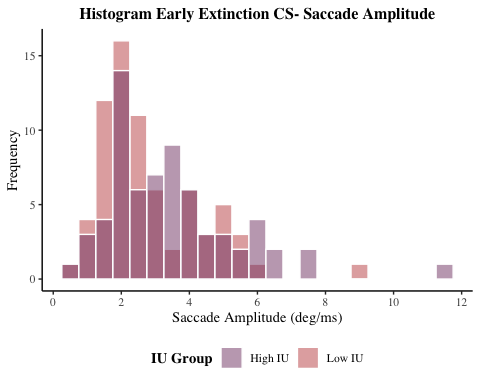
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_e\_ext\_csp\_sacc\_amplitude.png",   
 plot = hist\_e\_ext\_csp\_sacc\_amplitude,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

## Warning: Removed 1 rows containing non-finite values (stat\_bin).

### *Early Extinction CS-*

hist\_e\_ext\_csm\_sacc\_amplitude <- df %>%  
 ggplot(aes(e\_ext\_csm\_sacc\_amplitude, fill = iu\_group)) +  
 geom\_histogram(binwidth = .5, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 14, 2)) +  
 labs(x = "Saccade Amplitude (deg/ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Early Extinction CS- Saccade Amplitude") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_e\_ext\_csm\_sacc\_amplitude

## Warning: Removed 1 rows containing non-finite values (stat\_bin).



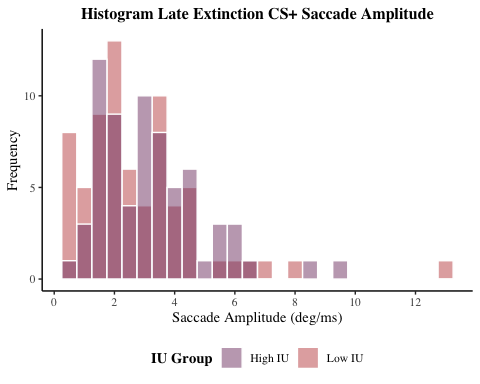
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_e\_ext\_csm\_sacc\_amplitude.png",   
 plot = hist\_e\_ext\_csm\_sacc\_amplitude,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

## Warning: Removed 1 rows containing non-finite values (stat\_bin).

### *Late Extinction CS+*

hist\_l\_ext\_csp\_sacc\_amplitude <- df %>%  
 ggplot(aes(l\_ext\_csp\_sacc\_amplitude, fill = iu\_group)) +  
 geom\_histogram(binwidth = .5, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 14, 2)) +  
 labs(x = "Saccade Amplitude (deg/ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Late Extinction CS+ Saccade Amplitude") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_l\_ext\_csp\_sacc\_amplitude

## Warning: Removed 1 rows containing non-finite values (stat\_bin).

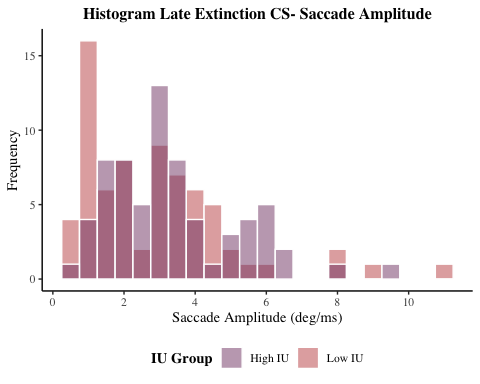


# save plot to file  
ggsave(filename = "graphs/histograms/hist\_l\_ext\_csp\_sacc\_amplitude.png",   
 plot = hist\_l\_ext\_csp\_sacc\_amplitude,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

## Warning: Removed 1 rows containing non-finite values (stat\_bin).

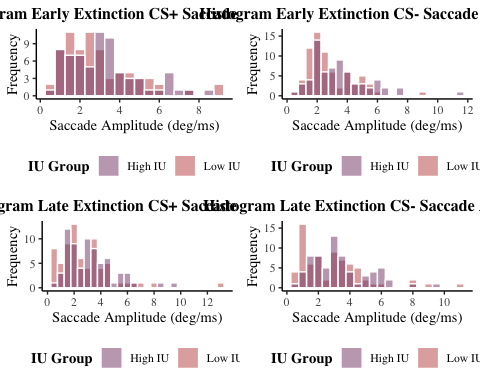
### *Late Extinction CS-*

hist\_l\_ext\_csm\_sacc\_amplitude <- df %>%  
 ggplot(aes(l\_ext\_csm\_sacc\_amplitude, fill = iu\_group)) +  
 geom\_histogram(binwidth = .5, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 14, 2)) +  
 labs(x = "Saccade Amplitude (deg/ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Late Extinction CS- Saccade Amplitude") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_l\_ext\_csm\_sacc\_amplitude



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_l\_ext\_csm\_sacc\_amplitude.png",   
 plot = hist\_l\_ext\_csm\_sacc\_amplitude,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
# combine extinction saccade amplitude graphs  
hists\_ext\_sacc\_amplitude <-   
 grid.arrange(hist\_e\_ext\_csp\_sacc\_amplitude, hist\_e\_ext\_csm\_sacc\_amplitude,  
 hist\_l\_ext\_csp\_sacc\_amplitude, hist\_l\_ext\_csm\_sacc\_amplitude,  
 ncol =2)

## Warning: Removed 1 rows containing non-finite values (stat\_bin).  
  
## Warning: Removed 1 rows containing non-finite values (stat\_bin).  
  
## Warning: Removed 1 rows containing non-finite values (stat\_bin).



# save plot to file  
ggsave(filename = "graphs/histograms/hists\_ext\_sacc\_amplitude.png",   
 plot = hists\_ext\_sacc\_amplitude,   
 width = 30,   
 height = 20,   
 dpi = 300,   
 units = "cm")

# *Descriptives*

## *Questionnaire Variables*

# for all participants  
descriptives\_all\_questionnaires <-   
 describe(df[, c("ius\_total", "sticsa\_total")], na.rm = TRUE)  
  
# for high IU group  
descriptives\_high\_iu\_questionnaires <-   
 describe(df[df$iu\_group == "1", c("ius\_total", "sticsa\_total")], na.rm = TRUE)  
  
# for low IU group  
descriptives\_low\_iu\_questionnaires <-   
 describe(df[df$iu\_group == "-1", c("ius\_total", "sticsa\_total")], na.rm = TRUE)  
  
# combine all into table  
descriptives\_questionnaires\_table <- round(rbind(descriptives\_all\_questionnaires,   
 descriptives\_high\_iu\_questionnaires,  
 descriptives\_low\_iu\_questionnaires), 2)  
  
# rename rows for easier interpretation  
rownames(descriptives\_questionnaires\_table) <- c("IUS 27 (All Participants)",  
 "STICSA Total (All Participants)",  
 "IUS 27 (High IU Group)",  
 "STICSA Total (High IU Group)",  
 "IUS 27 (Low IU Group)",  
 "STICSA Total (Low IU Group)")  
  
descriptives\_questionnaires\_table

## vars n mean sd median trimmed mad min  
## IUS 27 (All Participants) 1 139 65.82 20.39 63.0 64.27 20.76 32  
## STICSA Total (All Participants) 2 139 40.54 9.54 39.0 39.93 10.38 22  
## IUS 27 (High IU Group) 1 68 82.65 14.77 78.0 80.79 11.86 65  
## STICSA Total (High IU Group) 2 68 45.29 9.30 45.5 44.77 9.64 30  
## IUS 27 (Low IU Group) 1 71 49.70 8.51 51.0 49.96 10.38 32  
## STICSA Total (Low IU Group) 2 71 35.99 7.32 35.0 35.35 5.93 22  
## max range skew kurtosis se  
## IUS 27 (All Participants) 125 93 0.64 0.00 1.73  
## STICSA Total (All Participants) 69 47 0.65 0.06 0.81  
## IUS 27 (High IU Group) 125 60 1.11 0.53 1.79  
## STICSA Total (High IU Group) 69 39 0.47 -0.19 1.13  
## IUS 27 (Low IU Group) 64 32 -0.23 -1.05 1.01  
## STICSA Total (Low IU Group) 57 35 0.76 0.30 0.87

# write to csv  
write.csv(descriptives\_questionnaires\_table, file = "tables/descriptives/descriptives\_questionnaires\_table.csv",  
 row.names = TRUE)

## *Eye Movement Variables*

### *Fixation Count*

# for all participants  
descriptives\_all\_fix\_count <-   
 describe(df[, c("acq\_csp\_fix\_count","acq\_csm\_fix\_count",   
 "e\_ext\_csp\_fix\_count", "e\_ext\_csm\_fix\_count",  
 "l\_ext\_csp\_fix\_count", "l\_ext\_csm\_fix\_count")],   
 na.rm = TRUE)  
   
# for high IU group  
descriptives\_high\_iu\_fix\_count <-   
 describe(df[df$iu\_group == "1", c("acq\_csp\_fix\_count","acq\_csm\_fix\_count",   
 "e\_ext\_csp\_fix\_count", "e\_ext\_csm\_fix\_count",  
 "l\_ext\_csp\_fix\_count", "l\_ext\_csm\_fix\_count")],   
 na.rm = TRUE)  
   
# for low IU group  
descriptives\_low\_iu\_fix\_count <-   
 describe(df[df$iu\_group == "-1", c("acq\_csp\_fix\_count","acq\_csm\_fix\_count",   
 "e\_ext\_csp\_fix\_count", "e\_ext\_csm\_fix\_count",  
 "l\_ext\_csp\_fix\_count", "l\_ext\_csm\_fix\_count")],   
 na.rm = TRUE)  
  
# combine all into table  
descriptives\_fix\_count\_table <- round(rbind(descriptives\_all\_fix\_count,   
 descriptives\_high\_iu\_fix\_count,  
 descriptives\_low\_iu\_fix\_count), 2)  
  
# rename rows for easier interpretation  
rownames(descriptives\_fix\_count\_table) <- c("Acquisition CS+ Fix Count (All Participants)",   
 "Acquisition CS- Fix Count (All Participants)",   
 "Early Extinction CS+ Fix Count (All Participants)",   
 "Early Extinction CS- Fix Count (All Participants)",  
 "Late Extinction CS+ Fix Count (All Participants)",   
 "Late Extinction CS- Fix Count (All Participants)",  
 "Acquisition CS+ Fix Count (High IU Group)",   
 "Acquisition CS- Fix Count (High IU Group)",   
 "Early Extinction CS+ Fix Count (High IU Group)",   
 "Early Extinction CS- Fix Count (High IU Group)",  
 "Late Extinction CS+ Fix Count (High IU Group)",   
 "Late Extinction CS- Fix Count (High IU Group)",  
 "Acquisition CS+ Fix Count (Low IU Group)",   
 "Acquisition CS- Fix Count (Low IU Group)",   
 "Early Extinction CS+ Fix Count (Low IU Group)",   
 "Early Extinction CS- Fix Count (Low IU Group)",  
 "Late Extinction CS+ Fix Count (Low IU Group)",   
 "Late Extinction CS- Fix Count (Low IU Group)")  
  
descriptives\_fix\_count\_table

## vars n mean sd median  
## Acquisition CS+ Fix Count (All Participants) 1 139 6.90 3.65 6.67  
## Acquisition CS- Fix Count (All Participants) 2 139 7.31 3.25 6.75  
## Early Extinction CS+ Fix Count (All Participants) 3 139 7.16 3.70 6.50  
## Early Extinction CS- Fix Count (All Participants) 4 139 7.40 3.53 6.75  
## Late Extinction CS+ Fix Count (All Participants) 5 139 7.55 3.49 7.25  
## Late Extinction CS- Fix Count (All Participants) 6 139 7.86 3.52 7.75  
## Acquisition CS+ Fix Count (High IU Group) 1 68 7.51 3.84 7.08  
## Acquisition CS- Fix Count (High IU Group) 2 68 7.97 3.07 7.79  
## Early Extinction CS+ Fix Count (High IU Group) 3 68 7.54 3.26 6.75  
## Early Extinction CS- Fix Count (High IU Group) 4 68 8.14 3.26 7.88  
## Late Extinction CS+ Fix Count (High IU Group) 5 68 8.41 3.63 7.75  
## Late Extinction CS- Fix Count (High IU Group) 6 68 8.89 3.33 8.75  
## Acquisition CS+ Fix Count (Low IU Group) 1 71 6.33 3.38 5.50  
## Acquisition CS- Fix Count (Low IU Group) 2 71 6.67 3.31 5.92  
## Early Extinction CS+ Fix Count (Low IU Group) 3 71 6.80 4.06 6.00  
## Early Extinction CS- Fix Count (Low IU Group) 4 71 6.70 3.66 5.75  
## Late Extinction CS+ Fix Count (Low IU Group) 5 71 6.72 3.15 6.50  
## Late Extinction CS- Fix Count (Low IU Group) 6 71 6.87 3.43 6.50  
## trimmed mad min max range  
## Acquisition CS+ Fix Count (All Participants) 6.57 3.71 1.50 23.17 21.67  
## Acquisition CS- Fix Count (All Participants) 7.06 3.71 1.92 18.33 16.42  
## Early Extinction CS+ Fix Count (All Participants) 6.75 3.71 1.50 20.50 19.00  
## Early Extinction CS- Fix Count (All Participants) 7.14 3.71 1.50 21.50 20.00  
## Late Extinction CS+ Fix Count (All Participants) 7.33 3.34 1.00 22.00 21.00  
## Late Extinction CS- Fix Count (All Participants) 7.65 3.34 1.50 20.00 18.50  
## Acquisition CS+ Fix Count (High IU Group) 7.14 3.46 2.00 23.17 21.17  
## Acquisition CS- Fix Count (High IU Group) 7.78 3.21 2.92 18.33 15.42  
## Early Extinction CS+ Fix Count (High IU Group) 7.26 2.97 2.25 19.25 17.00  
## Early Extinction CS- Fix Count (High IU Group) 8.00 3.34 2.00 16.50 14.50  
## Late Extinction CS+ Fix Count (High IU Group) 8.11 3.71 1.50 22.00 20.50  
## Late Extinction CS- Fix Count (High IU Group) 8.61 2.41 3.25 20.00 16.75  
## Acquisition CS+ Fix Count (Low IU Group) 6.02 3.71 1.50 15.67 14.17  
## Acquisition CS- Fix Count (Low IU Group) 6.35 3.21 1.92 15.50 13.58  
## Early Extinction CS+ Fix Count (Low IU Group) 6.22 3.71 1.50 20.50 19.00  
## Early Extinction CS- Fix Count (Low IU Group) 6.31 3.34 1.50 21.50 20.00  
## Late Extinction CS+ Fix Count (Low IU Group) 6.57 3.34 1.00 16.00 15.00  
## Late Extinction CS- Fix Count (Low IU Group) 6.68 3.71 1.50 17.75 16.25  
## skew kurtosis se  
## Acquisition CS+ Fix Count (All Participants) 1.18 2.52 0.31  
## Acquisition CS- Fix Count (All Participants) 0.69 0.19 0.28  
## Early Extinction CS+ Fix Count (All Participants) 1.18 1.74 0.31  
## Early Extinction CS- Fix Count (All Participants) 0.82 0.96 0.30  
## Late Extinction CS+ Fix Count (All Participants) 0.83 1.34 0.30  
## Late Extinction CS- Fix Count (All Participants) 0.68 0.81 0.30  
## Acquisition CS+ Fix Count (High IU Group) 1.43 3.48 0.47  
## Acquisition CS- Fix Count (High IU Group) 0.74 0.76 0.37  
## Early Extinction CS+ Fix Count (High IU Group) 1.09 1.81 0.40  
## Early Extinction CS- Fix Count (High IU Group) 0.36 -0.42 0.39  
## Late Extinction CS+ Fix Count (High IU Group) 1.00 1.61 0.44  
## Late Extinction CS- Fix Count (High IU Group) 1.03 1.59 0.40  
## Acquisition CS+ Fix Count (Low IU Group) 0.75 -0.01 0.40  
## Acquisition CS- Fix Count (Low IU Group) 0.79 -0.13 0.39  
## Early Extinction CS+ Fix Count (Low IU Group) 1.30 1.65 0.48  
## Early Extinction CS- Fix Count (Low IU Group) 1.32 2.59 0.43  
## Late Extinction CS+ Fix Count (Low IU Group) 0.47 -0.15 0.37  
## Late Extinction CS- Fix Count (Low IU Group) 0.59 -0.05 0.41

# write to csv  
write.csv(descriptives\_fix\_count\_table, file = "tables/descriptives/descriptives\_fix\_count\_table.csv",  
 row.names = TRUE)

### *Fixation Duration*

# for all participants  
descriptives\_all\_fix\_duration <-   
 describe(df[, c("acq\_csp\_fix\_duration","acq\_csm\_fix\_duration",   
 "e\_ext\_csp\_fix\_duration", "e\_ext\_csm\_fix\_duration",  
 "l\_ext\_csp\_fix\_duration", "l\_ext\_csm\_fix\_duration")],   
 na.rm = TRUE)  
   
# for high IU group  
descriptives\_high\_iu\_fix\_duration <-   
 describe(df[df$iu\_group == "1", c("acq\_csp\_fix\_duration","acq\_csm\_fix\_duration",   
 "e\_ext\_csp\_fix\_duration", "e\_ext\_csm\_fix\_duration",   
 "l\_ext\_csp\_fix\_duration", "l\_ext\_csm\_fix\_duration")],   
 na.rm = TRUE)  
   
# for low IU group  
descriptives\_low\_iu\_fix\_duration <-   
 describe(df[df$iu\_group == "-1", c("acq\_csp\_fix\_duration","acq\_csm\_fix\_duration",   
 "e\_ext\_csp\_fix\_duration", "e\_ext\_csm\_fix\_duration",  
 "l\_ext\_csp\_fix\_duration", "l\_ext\_csm\_fix\_duration")],   
 na.rm = TRUE)  
  
# combine all in a table  
descriptives\_fix\_duration\_table <- round(rbind(descriptives\_all\_fix\_duration,   
 descriptives\_high\_iu\_fix\_duration,  
 descriptives\_low\_iu\_fix\_duration), 2)  
  
# rename rows for easier interpretation  
rownames(descriptives\_fix\_duration\_table) <- c("Acquisition CS+ Fix Duration (All Participants)",   
 "Acquisition CS- Fix Duration (All Participants)",   
 "Early Extinction CS+ Fix Duration (All Participants)",   
 "Early Extinction CS- Fix Duration (All Participants)",  
 "Late Extinction CS+ Fix Duration (All Participants)",   
 "Late Extinction CS- Fix Duration (All Participants)",  
 "Acquisition CS+ Fix Duration (High IU Group)",   
 "Acquisition CS- Fix Duration (High IU Group)",   
 "Early Extinction CS+ Fix Duration (High IU Group)",   
 "Early Extinction CS- Fix Duration (High IU Group)",  
 "Late Extinction CS+ Fix Duration (High IU Group)",   
 "Late Extinction CS- Fix Duration (High IU Group)",  
 "Acquisition CS+ Fix Duration (Low IU Group)",   
 "Acquisition CS- Fix Duration (Low IU Group)",   
 "Early Extinction CS+ Fix Duration (Low IU Group)",   
 "Early Extinction CS- Fix Duration (Low IU Group)",  
 "Late Extinction CS+ Fix Duration (Low IU Group)",   
 "Late Extinction CS- Fix Duration (Low IU Group)")  
  
descriptives\_fix\_duration\_table

## vars n mean sd  
## Acquisition CS+ Fix Duration (All Participants) 1 139 1309.36 1173.03  
## Acquisition CS- Fix Duration (All Participants) 2 139 1200.18 1048.80  
## Early Extinction CS+ Fix Duration (All Participants) 3 139 1104.04 930.02  
## Early Extinction CS- Fix Duration (All Participants) 4 139 1203.66 1288.87  
## Late Extinction CS+ Fix Duration (All Participants) 5 139 1066.13 1094.12  
## Late Extinction CS- Fix Duration (All Participants) 6 139 1068.60 1204.27  
## Acquisition CS+ Fix Duration (High IU Group) 1 68 1153.24 1126.41  
## Acquisition CS- Fix Duration (High IU Group) 2 68 1003.87 938.91  
## Early Extinction CS+ Fix Duration (High IU Group) 3 68 869.88 621.12  
## Early Extinction CS- Fix Duration (High IU Group) 4 68 833.27 912.11  
## Late Extinction CS+ Fix Duration (High IU Group) 5 68 799.03 732.10  
## Late Extinction CS- Fix Duration (High IU Group) 6 68 719.91 687.99  
## Acquisition CS+ Fix Duration (Low IU Group) 1 71 1458.89 1204.96  
## Acquisition CS- Fix Duration (Low IU Group) 2 71 1388.19 1118.70  
## Early Extinction CS+ Fix Duration (Low IU Group) 3 71 1328.31 1109.79  
## Early Extinction CS- Fix Duration (Low IU Group) 4 71 1558.41 1489.19  
## Late Extinction CS+ Fix Duration (Low IU Group) 5 71 1321.94 1308.18  
## Late Extinction CS- Fix Duration (Low IU Group) 6 71 1402.56 1474.73  
## median trimmed mad  
## Acquisition CS+ Fix Duration (All Participants) 789.44 1121.85 639.02  
## Acquisition CS- Fix Duration (All Participants) 778.02 1017.24 606.82  
## Early Extinction CS+ Fix Duration (All Participants) 786.25 958.21 611.53  
## Early Extinction CS- Fix Duration (All Participants) 674.06 937.16 519.93  
## Late Extinction CS+ Fix Duration (All Participants) 657.98 830.55 472.15  
## Late Extinction CS- Fix Duration (All Participants) 578.00 786.22 397.29  
## Acquisition CS+ Fix Duration (High IU Group) 666.40 943.03 510.01  
## Acquisition CS- Fix Duration (High IU Group) 649.90 830.74 412.10  
## Early Extinction CS+ Fix Duration (High IU Group) 716.65 780.80 516.83  
## Early Extinction CS- Fix Duration (High IU Group) 533.31 641.57 308.46  
## Late Extinction CS+ Fix Duration (High IU Group) 541.37 664.26 334.40  
## Late Extinction CS- Fix Duration (High IU Group) 510.20 585.21 306.25  
## Acquisition CS+ Fix Duration (Low IU Group) 1002.75 1309.89 888.32  
## Acquisition CS- Fix Duration (Low IU Group) 1081.07 1216.87 977.62  
## Early Extinction CS+ Fix Duration (Low IU Group) 931.86 1181.94 830.29  
## Early Extinction CS- Fix Duration (Low IU Group) 1017.70 1282.48 964.06  
## Late Extinction CS+ Fix Duration (Low IU Group) 781.46 1064.33 638.13  
## Late Extinction CS- Fix Duration (Low IU Group) 845.97 1102.87 689.98  
## min max range  
## Acquisition CS+ Fix Duration (All Participants) 87.39 5083.82 4996.43  
## Acquisition CS- Fix Duration (All Participants) 88.43 5446.78 5358.35  
## Early Extinction CS+ Fix Duration (All Participants) 79.01 5346.50 5267.49  
## Early Extinction CS- Fix Duration (All Participants) 65.23 6015.75 5950.52  
## Late Extinction CS+ Fix Duration (All Participants) 121.30 5923.00 5801.70  
## Late Extinction CS- Fix Duration (All Participants) 109.30 6086.56 5977.26  
## Acquisition CS+ Fix Duration (High IU Group) 87.39 5083.82 4996.43  
## Acquisition CS- Fix Duration (High IU Group) 88.43 5446.78 5358.35  
## Early Extinction CS+ Fix Duration (High IU Group) 110.04 3044.00 2933.96  
## Early Extinction CS- Fix Duration (High IU Group) 65.23 6015.75 5950.52  
## Late Extinction CS+ Fix Duration (High IU Group) 121.84 4252.33 4130.49  
## Late Extinction CS- Fix Duration (High IU Group) 109.30 4299.36 4190.06  
## Acquisition CS+ Fix Duration (Low IU Group) 129.50 4985.33 4855.84  
## Acquisition CS- Fix Duration (Low IU Group) 180.65 5219.17 5038.51  
## Early Extinction CS+ Fix Duration (Low IU Group) 79.01 5346.50 5267.49  
## Early Extinction CS- Fix Duration (Low IU Group) 119.97 5954.83 5834.86  
## Late Extinction CS+ Fix Duration (Low IU Group) 121.30 5923.00 5801.70  
## Late Extinction CS- Fix Duration (Low IU Group) 203.15 6086.56 5883.41  
## skew kurtosis se  
## Acquisition CS+ Fix Duration (All Participants) 1.41 1.29 99.50  
## Acquisition CS- Fix Duration (All Participants) 1.65 2.73 88.96  
## Early Extinction CS+ Fix Duration (All Participants) 1.58 2.76 78.88  
## Early Extinction CS- Fix Duration (All Participants) 2.05 3.83 109.32  
## Late Extinction CS+ Fix Duration (All Participants) 2.17 4.52 92.80  
## Late Extinction CS- Fix Duration (All Participants) 2.31 4.85 102.14  
## Acquisition CS+ Fix Duration (High IU Group) 1.94 3.43 136.60  
## Acquisition CS- Fix Duration (High IU Group) 2.30 6.41 113.86  
## Early Extinction CS+ Fix Duration (High IU Group) 1.50 2.47 75.32  
## Early Extinction CS- Fix Duration (High IU Group) 3.41 14.24 110.61  
## Late Extinction CS+ Fix Duration (High IU Group) 2.66 8.49 88.78  
## Late Extinction CS- Fix Duration (High IU Group) 3.05 10.94 83.43  
## Acquisition CS+ Fix Duration (Low IU Group) 0.98 -0.04 143.00  
## Acquisition CS- Fix Duration (Low IU Group) 1.21 1.01 132.77  
## Early Extinction CS+ Fix Duration (Low IU Group) 1.16 1.00 131.71  
## Early Extinction CS- Fix Duration (Low IU Group) 1.41 1.11 176.73  
## Late Extinction CS+ Fix Duration (Low IU Group) 1.64 1.90 155.25  
## Late Extinction CS- Fix Duration (Low IU Group) 1.64 1.62 175.02

# write to csv  
write.csv(descriptives\_fix\_duration\_table, file = "tables/descriptives/descriptives\_fix\_duration\_table.csv",   
 row.names = TRUE)

### *Saccade Amplitude*

# for all participants  
descriptives\_all\_sacc\_amplitude <-   
 describe(df[, c("acq\_csp\_sacc\_amplitude","acq\_csm\_sacc\_amplitude",   
 "e\_ext\_csp\_sacc\_amplitude", "e\_ext\_csm\_sacc\_amplitude",  
 "l\_ext\_csp\_sacc\_amplitude", "l\_ext\_csm\_sacc\_amplitude")],   
 na.rm = TRUE)  
   
# for high IU group  
descriptives\_high\_iu\_sacc\_amplitude <-   
 describe(df[df$iu\_group == "1", c("acq\_csp\_sacc\_amplitude","acq\_csm\_sacc\_amplitude",  
 "e\_ext\_csp\_sacc\_amplitude", "e\_ext\_csm\_sacc\_amplitude",  
 "l\_ext\_csp\_sacc\_amplitude", "l\_ext\_csm\_sacc\_amplitude")],   
 na.rm = TRUE)  
   
# for low IU group  
descriptives\_low\_iu\_sacc\_amplitude <-   
 describe(df[df$iu\_group == "-1", c("acq\_csp\_sacc\_amplitude","acq\_csm\_sacc\_amplitude",   
 "e\_ext\_csp\_sacc\_amplitude", "e\_ext\_csm\_sacc\_amplitude",  
 "l\_ext\_csp\_sacc\_amplitude", "l\_ext\_csm\_sacc\_amplitude")],   
 na.rm = TRUE)  
  
# combine all into one table  
descriptives\_sacc\_amplitude\_table <- round(rbind(descriptives\_all\_sacc\_amplitude,   
 descriptives\_high\_iu\_sacc\_amplitude,  
 descriptives\_low\_iu\_sacc\_amplitude), 2)  
  
# rename rows for easier interpretation  
rownames(descriptives\_sacc\_amplitude\_table) <- c("Acquisition CS+ Sacc Amplitude (All Participants)",   
 "Acquisition CS- Sacc Amplitude (All Participants)",   
 "Early Extinction CS+ Sacc Amplitude (All Participants)",   
 "Early Extinction CS- Sacc Amplitude (All Participants)",  
 "Late Extinction CS+ Sacc Amplitude (All Participants)",   
 "Late Extinction CS- Sacc Amplitude (All Participants)",  
 "Acquisition CS+ Sacc Amplitude (High IU Group)",   
 "Acquisition CS- Sacc Amplitude (High IU Group)",   
 "Early Extinction CS+ Sacc Amplitude (High IU Group)",   
 "Early Extinction CS- Sacc Amplitude (High IU Group)",  
 "Late Extinction CS+ Sacc Amplitude (High IU Group)",   
 "Late Extinction CS- Sacc Amplitude (High IU Group)",  
 "Acquisition CS+ Sacc Amplitude (Low IU Group)",   
 "Acquisition CS- Sacc Amplitude (Low IU Group)",   
 "Early Extinction CS+ Sacc Amplitude (Low IU Group)",   
 "Early Extinction CS- Sacc Amplitude (Low IU Group)",  
 "Late Extinction CS+ Sacc Amplitude (Low IU Group)",   
 "Late Extinction CS- Sacc Amplitude (Low IU Group)")  
  
descriptives\_sacc\_amplitude\_table

## vars n mean sd  
## Acquisition CS+ Sacc Amplitude (All Participants) 1 139 2.88 1.51  
## Acquisition CS- Sacc Amplitude (All Participants) 2 137 2.98 1.57  
## Early Extinction CS+ Sacc Amplitude (All Participants) 3 138 3.07 1.81  
## Early Extinction CS- Sacc Amplitude (All Participants) 4 138 3.13 1.73  
## Late Extinction CS+ Sacc Amplitude (All Participants) 5 138 3.00 1.92  
## Late Extinction CS- Sacc Amplitude (All Participants) 6 139 3.10 1.97  
## Acquisition CS+ Sacc Amplitude (High IU Group) 1 68 3.10 1.71  
## Acquisition CS- Sacc Amplitude (High IU Group) 2 67 3.16 1.70  
## Early Extinction CS+ Sacc Amplitude (High IU Group) 3 68 3.21 1.80  
## Early Extinction CS- Sacc Amplitude (High IU Group) 4 67 3.46 1.88  
## Late Extinction CS+ Sacc Amplitude (High IU Group) 5 68 3.21 1.78  
## Late Extinction CS- Sacc Amplitude (High IU Group) 6 68 3.37 1.85  
## Acquisition CS+ Sacc Amplitude (Low IU Group) 1 71 2.66 1.27  
## Acquisition CS- Sacc Amplitude (Low IU Group) 2 70 2.80 1.43  
## Early Extinction CS+ Sacc Amplitude (Low IU Group) 3 70 2.95 1.83  
## Early Extinction CS- Sacc Amplitude (Low IU Group) 4 71 2.81 1.53  
## Late Extinction CS+ Sacc Amplitude (Low IU Group) 5 70 2.79 2.03  
## Late Extinction CS- Sacc Amplitude (Low IU Group) 6 71 2.84 2.06  
## median trimmed mad min  
## Acquisition CS+ Sacc Amplitude (All Participants) 2.64 2.71 1.35 0.43  
## Acquisition CS- Sacc Amplitude (All Participants) 2.65 2.81 1.25 0.54  
## Early Extinction CS+ Sacc Amplitude (All Participants) 2.78 2.85 1.65 0.58  
## Early Extinction CS- Sacc Amplitude (All Participants) 2.66 2.94 1.35 0.42  
## Late Extinction CS+ Sacc Amplitude (All Participants) 2.69 2.78 1.61 0.38  
## Late Extinction CS- Sacc Amplitude (All Participants) 2.90 2.86 1.89 0.42  
## Acquisition CS+ Sacc Amplitude (High IU Group) 2.99 2.92 1.74 0.43  
## Acquisition CS- Sacc Amplitude (High IU Group) 2.86 2.96 1.49 0.54  
## Early Extinction CS+ Sacc Amplitude (High IU Group) 3.08 3.02 1.75 0.64  
## Early Extinction CS- Sacc Amplitude (High IU Group) 3.18 3.28 1.74 0.69  
## Late Extinction CS+ Sacc Amplitude (High IU Group) 2.90 3.03 1.78 0.38  
## Late Extinction CS- Sacc Amplitude (High IU Group) 3.13 3.22 1.90 0.61  
## Acquisition CS+ Sacc Amplitude (Low IU Group) 2.52 2.56 1.17 0.59  
## Acquisition CS- Sacc Amplitude (Low IU Group) 2.60 2.66 1.23 0.59  
## Early Extinction CS+ Sacc Amplitude (Low IU Group) 2.48 2.70 1.52 0.58  
## Early Extinction CS- Sacc Amplitude (Low IU Group) 2.34 2.65 1.07 0.42  
## Late Extinction CS+ Sacc Amplitude (Low IU Group) 2.25 2.52 1.69 0.43  
## Late Extinction CS- Sacc Amplitude (Low IU Group) 2.63 2.53 1.93 0.42  
## max range skew  
## Acquisition CS+ Sacc Amplitude (All Participants) 8.15 7.72 1.01  
## Acquisition CS- Sacc Amplitude (All Participants) 8.57 8.04 1.12  
## Early Extinction CS+ Sacc Amplitude (All Participants) 9.18 8.59 1.14  
## Early Extinction CS- Sacc Amplitude (All Participants) 11.42 11.00 1.44  
## Late Extinction CS+ Sacc Amplitude (All Participants) 13.11 12.73 1.72  
## Late Extinction CS- Sacc Amplitude (All Participants) 10.95 10.53 1.25  
## Acquisition CS+ Sacc Amplitude (High IU Group) 8.15 7.72 0.96  
## Acquisition CS- Sacc Amplitude (High IU Group) 8.57 8.04 1.14  
## Early Extinction CS+ Sacc Amplitude (High IU Group) 8.65 8.01 0.89  
## Early Extinction CS- Sacc Amplitude (High IU Group) 11.42 10.73 1.38  
## Late Extinction CS+ Sacc Amplitude (High IU Group) 9.62 9.24 1.11  
## Late Extinction CS- Sacc Amplitude (High IU Group) 9.74 9.13 0.93  
## Acquisition CS+ Sacc Amplitude (Low IU Group) 6.35 5.76 0.68  
## Acquisition CS- Sacc Amplitude (Low IU Group) 7.37 6.78 0.93  
## Early Extinction CS+ Sacc Amplitude (Low IU Group) 9.18 8.59 1.37  
## Early Extinction CS- Sacc Amplitude (Low IU Group) 9.11 8.69 1.33  
## Late Extinction CS+ Sacc Amplitude (Low IU Group) 13.11 12.68 2.20  
## Late Extinction CS- Sacc Amplitude (Low IU Group) 10.95 10.53 1.57  
## kurtosis se  
## Acquisition CS+ Sacc Amplitude (All Participants) 0.86 0.13  
## Acquisition CS- Sacc Amplitude (All Participants) 1.35 0.13  
## Early Extinction CS+ Sacc Amplitude (All Participants) 1.13 0.15  
## Early Extinction CS- Sacc Amplitude (All Participants) 3.30 0.15  
## Late Extinction CS+ Sacc Amplitude (All Participants) 5.31 0.16  
## Late Extinction CS- Sacc Amplitude (All Participants) 1.95 0.17  
## Acquisition CS+ Sacc Amplitude (High IU Group) 0.40 0.21  
## Acquisition CS- Sacc Amplitude (High IU Group) 1.13 0.21  
## Early Extinction CS+ Sacc Amplitude (High IU Group) 0.32 0.22  
## Early Extinction CS- Sacc Amplitude (High IU Group) 3.07 0.23  
## Late Extinction CS+ Sacc Amplitude (High IU Group) 1.55 0.22  
## Late Extinction CS- Sacc Amplitude (High IU Group) 0.76 0.22  
## Acquisition CS+ Sacc Amplitude (Low IU Group) -0.16 0.15  
## Acquisition CS- Sacc Amplitude (Low IU Group) 0.81 0.17  
## Early Extinction CS+ Sacc Amplitude (Low IU Group) 1.92 0.22  
## Early Extinction CS- Sacc Amplitude (Low IU Group) 2.40 0.18  
## Late Extinction CS+ Sacc Amplitude (Low IU Group) 7.87 0.24  
## Late Extinction CS- Sacc Amplitude (Low IU Group) 3.05 0.25

# write to csv  
write.csv(descriptives\_sacc\_amplitude\_table, file = "tables/descriptives/descriptives\_sacc\_amplitude\_table.csv",   
 row.names = TRUE)

# *Data Transformation*

## *Log-Transformation of Fixation Duration*

# as fixation duration had high skew (>3) in high IU group for early and late   
# extinction CS-, fixation duration will be log-transformed for each condition  
  
# for acquisition CS+  
df$acq\_csp\_fix\_duration\_log <- log(df$acq\_csp\_fix\_duration)  
  
# for acquisition CS-  
df$acq\_csm\_fix\_duration\_log <- log(df$acq\_csm\_fix\_duration)  
  
# for early extinction CS+  
df$e\_ext\_csp\_fix\_duration\_log <- log(df$e\_ext\_csp\_fix\_duration)  
  
# for early extinction CS-  
df$e\_ext\_csm\_fix\_duration\_log <- log(df$e\_ext\_csm\_fix\_duration)  
  
# for late extinction CS+  
df$l\_ext\_csp\_fix\_duration\_log <- log(df$l\_ext\_csp\_fix\_duration)  
  
# for late extinction CS-  
df$l\_ext\_csm\_fix\_duration\_log <- log(df$l\_ext\_csm\_fix\_duration)

## *Check Descriptives of Fixation Duration Following Log-Transformation*

# re-compute descriptives for fixation duration following log transformation  
  
# for all participants  
descriptives\_all\_fix\_duration\_log <-   
 describe(df[, c("acq\_csp\_fix\_duration\_log","acq\_csm\_fix\_duration\_log",   
 "e\_ext\_csp\_fix\_duration\_log", "e\_ext\_csm\_fix\_duration\_log",  
 "l\_ext\_csp\_fix\_duration\_log", "l\_ext\_csm\_fix\_duration\_log")],   
 na.rm = TRUE)  
   
# for high IU group  
descriptives\_high\_iu\_fix\_duration\_log <-   
 describe(df[df$iu\_group == "1", c("acq\_csp\_fix\_duration\_log","acq\_csm\_fix\_duration\_log",   
 "e\_ext\_csp\_fix\_duration\_log", "e\_ext\_csm\_fix\_duration\_log",   
 "l\_ext\_csp\_fix\_duration\_log", "l\_ext\_csm\_fix\_duration\_log")],   
 na.rm = TRUE)  
   
# for low IU group  
descriptives\_low\_iu\_fix\_duration\_log <-   
 describe(df[df$iu\_group == "-1", c("acq\_csp\_fix\_duration\_log","acq\_csm\_fix\_duration\_log",   
 "e\_ext\_csp\_fix\_duration\_log", "e\_ext\_csm\_fix\_duration\_log",  
 "l\_ext\_csp\_fix\_duration\_log", "l\_ext\_csm\_fix\_duration\_log")],   
 na.rm = TRUE)  
  
# combine all to table  
descriptives\_fix\_duration\_table\_log <- round(rbind(descriptives\_all\_fix\_duration\_log,   
 descriptives\_high\_iu\_fix\_duration\_log,  
 descriptives\_low\_iu\_fix\_duration\_log), 2)  
  
# rename rows for easier interpretation  
rownames(descriptives\_fix\_duration\_table\_log) <- c("Acquisition CS+ Fix Duration (All Participants)",   
 "Acquisition CS- Fix Duration (All Participants)",   
 "Early Extinction CS+ Fix Duration (All Participants)",   
 "Early Extinction CS- Fix Duration (All Participants)",  
 "Late Extinction CS+ Fix Duration (All Participants)",   
 "Late Extinction CS- Fix Duration (All Participants)",  
 "Acquisition CS+ Fix Duration (High IU Group)",   
 "Acquisition CS- Fix Duration (High IU Group)",   
 "Early Extinction CS+ Fix Duration (High IU Group)",   
 "Early Extinction CS- Fix Duration (High IU Group)",  
 "Late Extinction CS+ Fix Duration (High IU Group)",   
 "Late Extinction CS- Fix Duration (High IU Group)",  
 "Acquisition CS+ Fix Duration (Low IU Group)",   
 "Acquisition CS- Fix Duration (Low IU Group)",   
 "Early Extinction CS+ Fix Duration (Low IU Group)",   
 "Early Extinction CS- Fix Duration (Low IU Group)",  
 "Late Extinction CS+ Fix Duration (Low IU Group)",   
 "Late Extinction CS- Fix Duration (Low IU Group)")  
  
descriptives\_fix\_duration\_table\_log

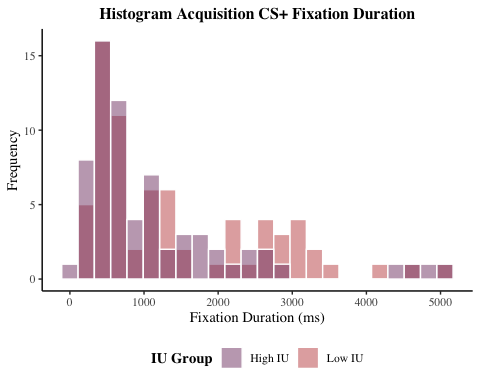
## vars n mean sd median  
## Acquisition CS+ Fix Duration (All Participants) 1 139 6.80 0.89 6.67  
## Acquisition CS- Fix Duration (All Participants) 2 139 6.76 0.83 6.66  
## Early Extinction CS+ Fix Duration (All Participants) 3 139 6.68 0.84 6.67  
## Early Extinction CS- Fix Duration (All Participants) 4 139 6.67 0.90 6.51  
## Late Extinction CS+ Fix Duration (All Participants) 5 139 6.60 0.83 6.49  
## Late Extinction CS- Fix Duration (All Participants) 6 139 6.56 0.85 6.36  
## Acquisition CS+ Fix Duration (High IU Group) 1 68 6.68 0.87 6.50  
## Acquisition CS- Fix Duration (High IU Group) 2 68 6.60 0.78 6.48  
## Early Extinction CS+ Fix Duration (High IU Group) 3 68 6.54 0.70 6.57  
## Early Extinction CS- Fix Duration (High IU Group) 4 68 6.40 0.75 6.28  
## Late Extinction CS+ Fix Duration (High IU Group) 5 68 6.41 0.72 6.29  
## Late Extinction CS- Fix Duration (High IU Group) 6 68 6.31 0.71 6.23  
## Acquisition CS+ Fix Duration (Low IU Group) 1 71 6.92 0.89 6.91  
## Acquisition CS- Fix Duration (Low IU Group) 2 71 6.91 0.85 6.99  
## Early Extinction CS+ Fix Duration (Low IU Group) 3 71 6.81 0.94 6.84  
## Early Extinction CS- Fix Duration (Low IU Group) 4 71 6.92 0.96 6.93  
## Late Extinction CS+ Fix Duration (Low IU Group) 5 71 6.79 0.89 6.66  
## Late Extinction CS- Fix Duration (Low IU Group) 6 71 6.81 0.91 6.74  
## trimmed mad min max  
## Acquisition CS+ Fix Duration (All Participants) 6.80 0.94 4.47 8.53  
## Acquisition CS- Fix Duration (All Participants) 6.74 0.97 4.48 8.60  
## Early Extinction CS+ Fix Duration (All Participants) 6.69 0.93 4.37 8.58  
## Early Extinction CS- Fix Duration (All Participants) 6.63 0.84 4.18 8.70  
## Late Extinction CS+ Fix Duration (All Participants) 6.55 0.86 4.80 8.69  
## Late Extinction CS- Fix Duration (All Participants) 6.50 0.71 4.69 8.71  
## Acquisition CS+ Fix Duration (High IU Group) 6.67 0.79 4.47 8.53  
## Acquisition CS- Fix Duration (High IU Group) 6.57 0.74 4.48 8.60  
## Early Extinction CS+ Fix Duration (High IU Group) 6.54 0.77 4.70 8.02  
## Early Extinction CS- Fix Duration (High IU Group) 6.35 0.67 4.18 8.70  
## Late Extinction CS+ Fix Duration (High IU Group) 6.38 0.75 4.80 8.36  
## Late Extinction CS- Fix Duration (High IU Group) 6.29 0.61 4.69 8.37  
## Acquisition CS+ Fix Duration (Low IU Group) 6.94 1.15 4.86 8.51  
## Acquisition CS- Fix Duration (Low IU Group) 6.91 1.01 5.20 8.56  
## Early Extinction CS+ Fix Duration (Low IU Group) 6.86 1.15 4.37 8.58  
## Early Extinction CS- Fix Duration (Low IU Group) 6.91 1.14 4.79 8.69  
## Late Extinction CS+ Fix Duration (Low IU Group) 6.75 0.91 4.80 8.69  
## Late Extinction CS- Fix Duration (Low IU Group) 6.75 0.96 5.31 8.71  
## range skew kurtosis se  
## Acquisition CS+ Fix Duration (All Participants) 4.06 0.02 -0.64 0.08  
## Acquisition CS- Fix Duration (All Participants) 4.12 0.11 -0.65 0.07  
## Early Extinction CS+ Fix Duration (All Participants) 4.21 -0.11 -0.40 0.07  
## Early Extinction CS- Fix Duration (All Participants) 4.52 0.34 -0.31 0.08  
## Late Extinction CS+ Fix Duration (All Participants) 3.89 0.45 -0.27 0.07  
## Late Extinction CS- Fix Duration (All Participants) 4.02 0.59 -0.02 0.07  
## Acquisition CS+ Fix Duration (High IU Group) 4.06 0.09 -0.21 0.11  
## Acquisition CS- Fix Duration (High IU Group) 4.12 0.25 -0.01 0.09  
## Early Extinction CS+ Fix Duration (High IU Group) 3.32 -0.08 -0.48 0.08  
## Early Extinction CS- Fix Duration (High IU Group) 4.52 0.48 1.07 0.09  
## Late Extinction CS+ Fix Duration (High IU Group) 3.55 0.41 -0.03 0.09  
## Late Extinction CS- Fix Duration (High IU Group) 3.67 0.35 0.66 0.09  
## Acquisition CS+ Fix Duration (Low IU Group) 3.65 -0.07 -1.03 0.11  
## Acquisition CS- Fix Duration (Low IU Group) 3.36 -0.07 -1.05 0.10  
## Early Extinction CS+ Fix Duration (Low IU Group) 4.21 -0.32 -0.53 0.11  
## Early Extinction CS- Fix Duration (Low IU Group) 3.90 0.03 -0.87 0.11  
## Late Extinction CS+ Fix Duration (Low IU Group) 3.89 0.29 -0.67 0.11  
## Late Extinction CS- Fix Duration (Low IU Group) 3.40 0.47 -0.83 0.11

# write to csv  
write.csv(descriptives\_fix\_duration\_table\_log, file = "tables/descriptives/descriptives\_fix\_duration\_table\_log.csv",   
 row.names = TRUE)  
  
### there are no longer any skew values of +/- 3.

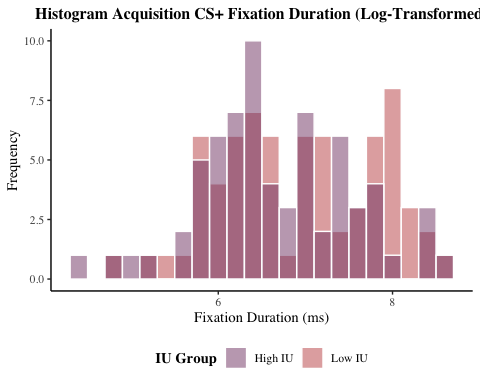
## *Check Histograms of Fixation Duration Following Log-Transformation*

### *Acquisition CS+*

########## pre-log-transformation  
hist\_acq\_csp\_fix\_duration



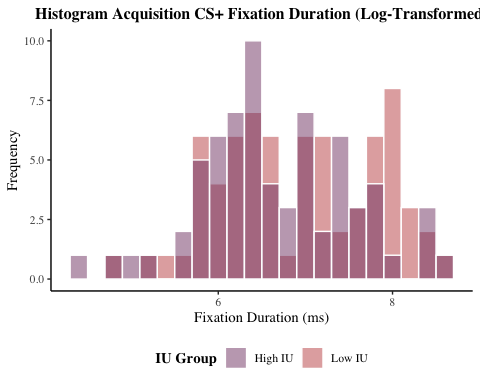
########## post-log-transformation  
hist\_acq\_csp\_fix\_duration\_log <- df %>%  
 ggplot(aes(acq\_csp\_fix\_duration\_log, fill = iu\_group)) +  
 geom\_histogram(binwidth = .2, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 12, 2)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Acquisition CS+ Fixation Duration (Log-Transformed)") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_acq\_csp\_fix\_duration\_log



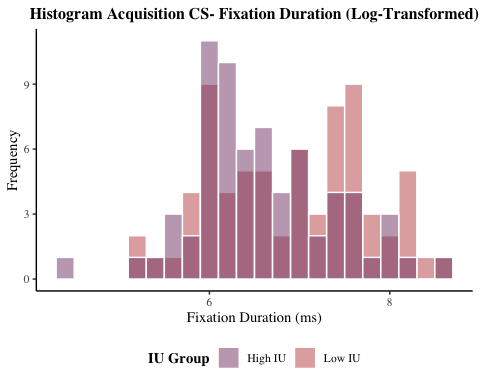
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_acq\_csp\_fix\_duration\_log.png",   
 plot = hist\_acq\_csp\_fix\_duration\_log,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Acquisition CS-*

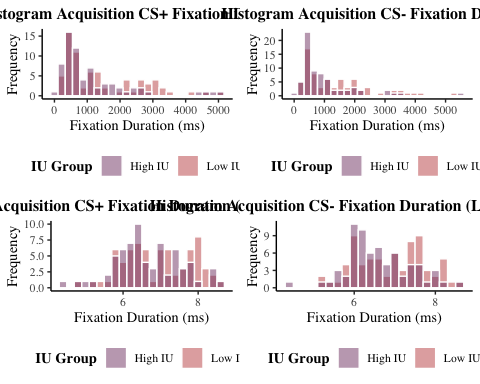
########## pre-log-transformation  
hist\_acq\_csp\_fix\_duration\_log



########## post-log-transformation  
hist\_acq\_csm\_fix\_duration\_log <- df %>%  
 ggplot(aes(acq\_csm\_fix\_duration\_log, fill = iu\_group)) +  
 geom\_histogram(binwidth = .2, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 12, 2)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Acquisition CS- Fixation Duration (Log-Transformed)") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_acq\_csm\_fix\_duration\_log



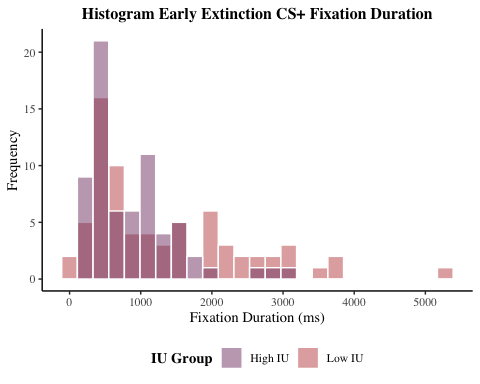
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_acq\_csm\_fix\_duration\_log.png",   
 plot = hist\_acq\_csm\_fix\_duration\_log,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
# combine histograms of acquisition fix duration pre and post log-transformation  
hists\_fix\_duration\_acq\_log <- grid.arrange(hist\_acq\_csp\_fix\_duration, hist\_acq\_csm\_fix\_duration,  
 hist\_acq\_csp\_fix\_duration\_log, hist\_acq\_csm\_fix\_duration\_log,  
 ncol = 2)



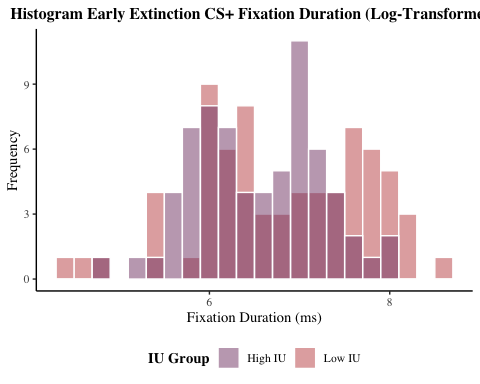
# save plot to file  
ggsave(filename = "graphs/histograms/hists\_fix\_duration\_acq\_log.png",   
 plot = hists\_fix\_duration\_acq\_log,   
 width = 30,   
 height = 20,   
 dpi = 300,   
 units = "cm")

### *Early Extinction CS+*

########## pre-log-transformation  
hist\_e\_ext\_csp\_fix\_duration



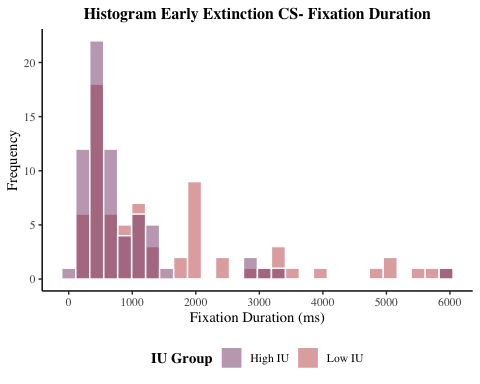
########## post-log-transformation  
hist\_e\_ext\_csp\_fix\_duration\_log <- df %>%  
 ggplot(aes(e\_ext\_csp\_fix\_duration\_log, fill = iu\_group)) +  
 geom\_histogram(binwidth = .2, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 12, 2)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Early Extinction CS+ Fixation Duration (Log-Transformed)") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_e\_ext\_csp\_fix\_duration\_log



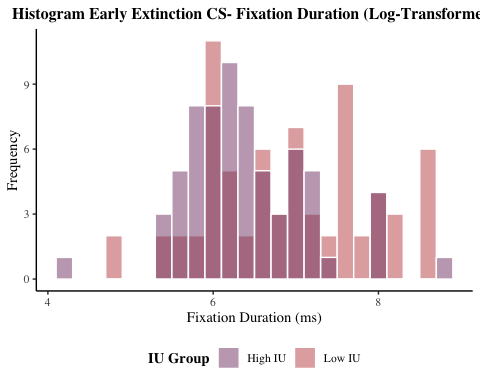
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_e\_ext\_csp\_fix\_duration\_log.png",   
 plot = hist\_e\_ext\_csp\_fix\_duration\_log,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Early Extinction CS-*

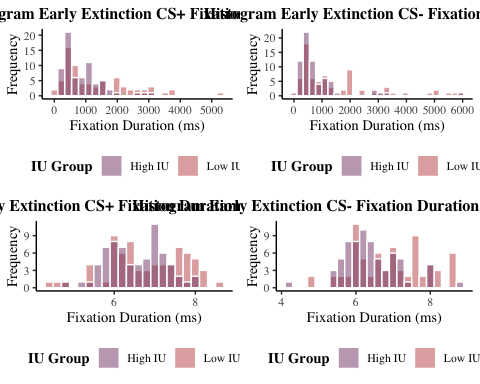
########## pre-log-transformation  
hist\_e\_ext\_csm\_fix\_duration



########## post-log-transformation  
hist\_e\_ext\_csm\_fix\_duration\_log <- df %>%  
 ggplot(aes(e\_ext\_csm\_fix\_duration\_log, fill = iu\_group)) +  
 geom\_histogram(binwidth = .2, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 12, 2)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Early Extinction CS- Fixation Duration (Log-Transformed)") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_e\_ext\_csm\_fix\_duration\_log



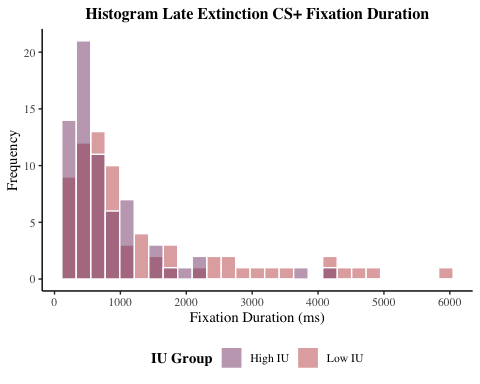
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_e\_ext\_csm\_fix\_duration\_log.png",   
 plot = hist\_e\_ext\_csm\_fix\_duration\_log,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
# combine histograms of early extinction fix duration pre and post log-transformation  
hists\_fix\_duration\_e\_ext\_log <- grid.arrange(hist\_e\_ext\_csp\_fix\_duration, hist\_e\_ext\_csm\_fix\_duration,  
 hist\_e\_ext\_csp\_fix\_duration\_log, hist\_e\_ext\_csm\_fix\_duration\_log,  
 ncol = 2)



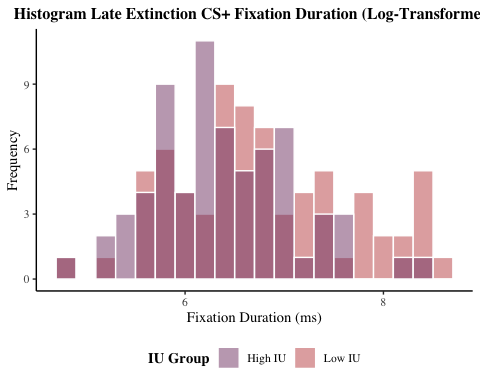
# save plot to file  
ggsave(filename = "graphs/histograms/hists\_fix\_duration\_e\_ext\_log.png",   
 plot = hists\_fix\_duration\_e\_ext\_log,   
 width = 30,   
 height = 20,   
 dpi = 300,   
 units = "cm")

### *Late Extinction CS+*

########## pre-log-transformation  
hist\_l\_ext\_csp\_fix\_duration



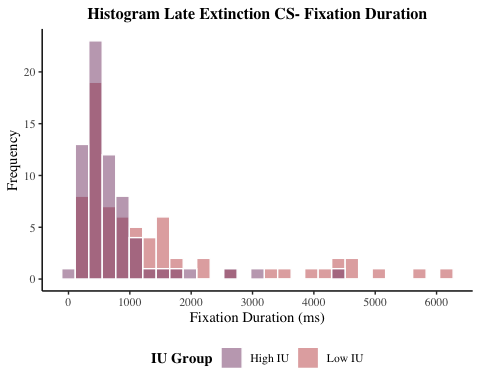
########## post-log-transformation  
hist\_l\_ext\_csp\_fix\_duration\_log <- df %>%  
 ggplot(aes(l\_ext\_csp\_fix\_duration\_log, fill = iu\_group)) +  
 geom\_histogram(binwidth = .2, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 12, 2)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Late Extinction CS+ Fixation Duration (Log-Transformed)") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_l\_ext\_csp\_fix\_duration\_log



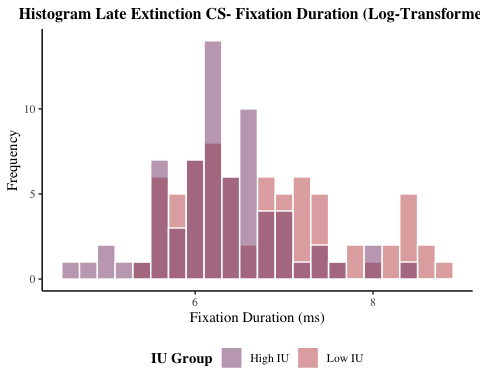
# save plot to file  
ggsave(filename = "graphs/histograms/hist\_l\_ext\_csp\_fix\_duration\_log.png",   
 plot = hist\_l\_ext\_csp\_fix\_duration\_log,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

### *Late Extinction CS-*

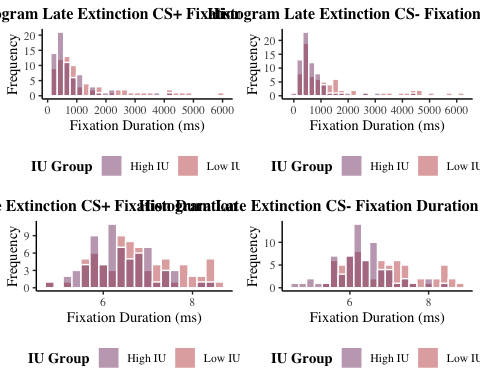
########## pre-log-transformation  
hist\_l\_ext\_csm\_fix\_duration



########## post-log-transformation  
hist\_l\_ext\_csm\_fix\_duration\_log <- df %>%  
 ggplot(aes(l\_ext\_csm\_fix\_duration\_log, fill = iu\_group)) +  
 geom\_histogram(binwidth = .2, colour = "white", alpha = .5, position = "identity") +  
 theme\_classic() +  
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_text(face = "bold", hjust = 0.5, size = 15)) +  
 scale\_x\_continuous(breaks = seq(0, 12, 2)) +  
 labs(x = "Fixation Duration (ms)", y = "Frequency") +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 ggtitle("Histogram Late Extinction CS- Fixation Duration (Log-Transformed)") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold")) +  
 guides(fill = guide\_legend(reverse = TRUE)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(fill = "IU Group")   
hist\_l\_ext\_csm\_fix\_duration\_log



# save plot to file  
ggsave(filename = "graphs/histograms/hist\_l\_ext\_csm\_fix\_duration\_log.png",   
 plot = hist\_l\_ext\_csm\_fix\_duration\_log,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")  
  
# combine histograms of late extinction fix duration pre and post log-transformation  
hists\_fix\_duration\_l\_ext\_log <- grid.arrange(hist\_l\_ext\_csp\_fix\_duration, hist\_l\_ext\_csm\_fix\_duration,  
 hist\_l\_ext\_csp\_fix\_duration\_log, hist\_l\_ext\_csm\_fix\_duration\_log,  
 ncol = 2)



# save plot to file  
ggsave(filename = "graphs/histograms/hists\_fix\_duration\_l\_ext\_log.png",   
 plot = hists\_fix\_duration\_l\_ext\_log,   
 width = 30,   
 height = 20,   
 dpi = 300,   
 units = "cm")

# *ANOVAs*

## *ANOVA Acquisition Fixation Count*

# transform wide format data into long format for mixed ANOVA   
df\_long\_acq\_fix\_count <- melt(df, id = c("id", "iu\_group"),   
 measure.vars = c("acq\_csp\_fix\_count",   
 "acq\_csm\_fix\_count"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_acq\_fix\_count) = c("id", "iu\_group", "condition", "fix\_count")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_acq\_fix\_count$stimulus <-   
 factor(ifelse(df\_long\_acq\_fix\_count$condition == "acq\_csp\_fix\_count", 1, -1))  
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) mixed ANOVA,   
# and obtain effect size (partial eta squared)  
acq\_fix\_count\_anova <-   
 anova\_test(df\_long\_acq\_fix\_count, fix\_count ~ iu\_group \* stimulus + Error(id/stimulus),  
 effect.size = "pes")   
  
# obtain the mixed ANOVA results  
get\_anova\_table(acq\_fix\_count\_anova)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 iu\_group 1 137 4.806 0.030000 \* 0.034  
## 2 stimulus 1 137 11.441 0.000937 \* 0.077  
## 3 iu\_group:stimulus 1 137 0.258 0.613000 0.002

# results:  
# IU: F(1,137) = 4.81, p = .030\*, eta2(partial) = .034  
# Stimulus: F(1,137) = 11.44, p < .001\*\*\*, eta2(partial) = .077  
# IU \* Stimulus: F(1, 137) = 0.26, p = .613, eta2(partial) = .002  
  
# therefore, there is a significant effect of IU & Stimulus on fixation count in acquisition,   
# and no significant IU\*Stimulus interaction  
  
# write to csv  
write.csv((get\_anova\_table(acq\_fix\_count\_anova)),   
 file = "tables/anovas/acq\_fix\_count\_anova.csv")

## *ANOVA Acquisition Fixation Duration (Log Transformed)*

# transform wide format data into long format for mixed ANOVA   
df\_long\_acq\_fix\_duration\_log <- melt(df, id = c("id", "iu\_group"),   
 measure.vars = c("acq\_csp\_fix\_duration\_log",   
 "acq\_csm\_fix\_duration\_log"))  
  
# the first input for melt command is the df with wide format. Second input  
# is id =, which is where we list ppts with specific variables within wide format  
# df. Here we have ppts ID no's as participant specific variable, and IU   
# group they are assigned to is also specific for each participant. Therefore  
# only need to list the two variables after id =.   
  
# rename columns for easier interpretation  
colnames(df\_long\_acq\_fix\_duration\_log) = c("id", "iu\_group", "condition", "fix\_duration\_log")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_acq\_fix\_duration\_log$stimulus <-   
 factor(ifelse(df\_long\_acq\_fix\_duration\_log$condition == "acq\_csp\_fix\_duration\_log", 1, -1))  
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) mixed ANOVA,   
# and obtain effect size (partial eta squared)  
acq\_fix\_duration\_anova\_log <-   
 anova\_test(df\_long\_acq\_fix\_duration\_log, fix\_duration\_log ~ iu\_group \* stimulus + Error(id/stimulus),  
 effect.size = "pes")   
# the error(id/stimulus) variable is unique to repeated-measures ANOVA, and means  
# that the variable 'stimulus' is manipulated within 'id'  
  
# obtain the mixed ANOVA results  
get\_anova\_table(acq\_fix\_duration\_anova\_log)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 iu\_group 1 137 3.907 0.050 0.028  
## 2 stimulus 1 137 2.921 0.090 0.021  
## 3 iu\_group:stimulus 1 137 1.271 0.261 0.009

# results:  
# IU: F(1,137) = 3.91, p = .050\*, eta2(partial) = .028  
# Stimulus: F(1,137) = 2.92, p = .090, eta2(partial) = .021  
# IU \* Stimulus: F(1, 137) = 1.27, p = .261, eta2(partial) = .009  
  
# therefore, there is a sig effect of IU, and no  
# sig effect of stimulus or IU-stimulus interaction   
  
# write to csv  
write.csv((get\_anova\_table(acq\_fix\_duration\_anova\_log)),   
 file = "tables/anovas/acq\_fix\_duration\_anova\_log.csv")

## *ANOVA Acquisition Saccade Amplitude*

# transform wide format data into long format for mixed ANOVA   
df\_long\_acq\_sacc\_amplitude <- melt(df, id = c("id", "iu\_group"),   
 measure.vars = c("acq\_csp\_sacc\_amplitude",   
 "acq\_csm\_sacc\_amplitude"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_acq\_sacc\_amplitude) = c("id", "iu\_group", "condition", "sacc\_amplitude")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_acq\_sacc\_amplitude$stimulus <-   
 factor(ifelse(df\_long\_acq\_sacc\_amplitude$condition == "acq\_csp\_sacc\_amplitude", 1, -1))  
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) mixed ANOVA,   
# and obtain effect size (partial eta squared)  
acq\_sacc\_amplitude\_anova <-   
 anova\_test(df\_long\_acq\_sacc\_amplitude, sacc\_amplitude ~ iu\_group \* stimulus + Error(id/stimulus),  
 effect.size = "pes")

## Warning: NA detected in rows: 234,259.  
## Removing this rows before the analysis.

# obtain the mixed ANOVA results  
get\_anova\_table(acq\_sacc\_amplitude\_anova)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 iu\_group 1 135 2.984 0.086 0.022  
## 2 stimulus 1 135 0.950 0.332 0.007  
## 3 iu\_group:stimulus 1 135 0.379 0.539 0.003

# results:  
# IU: F(1,135) = 2.98, p = .086, eta2(partial) = .022  
# Stimulus: F(1,135) = 0.95, p = .332, eta2(partial) = .007  
# IU \* Stimulus: F(1, 135) = 0.38, p = .539, eta2(partial) = .003  
  
# therefore, there are no significant effects on saccade amplitude in  
# acquisition  
  
# write to csv  
write.csv((get\_anova\_table(acq\_sacc\_amplitude\_anova)),   
 file = "tables/anovas/acq\_sacc\_amplitude\_anova.csv")

## *ANOVA Extinction Fixation Count*

# transform wide format data into long format for mixed ANOVA   
df\_long\_ext\_fix\_count <- melt(df, id = c("id", "iu\_group"),   
 measure.vars = c("e\_ext\_csp\_fix\_count",   
 "e\_ext\_csm\_fix\_count",  
 "l\_ext\_csp\_fix\_count",   
 "l\_ext\_csm\_fix\_count"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_ext\_fix\_count) = c("id", "iu\_group", "condition", "fix\_count")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_ext\_fix\_count$stimulus <-   
 factor(ifelse(df\_long\_ext\_fix\_count$condition == "e\_ext\_csp\_fix\_count" |   
 df\_long\_ext\_fix\_count$condition == "l\_ext\_csp\_fix\_count", 1, -1))  
  
# create column to code extinction as early (1) and late (-1)  
df\_long\_ext\_fix\_count$time <-   
 factor(ifelse(df\_long\_ext\_fix\_count$condition == "e\_ext\_csp\_fix\_count" |   
 df\_long\_ext\_fix\_count$condition == "e\_ext\_csm\_fix\_count", 1, -1))  
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) x 2 (Time: Early, Late) mixed ANOVA,   
# and obtain effect size (partial eta squared)  
ext\_fix\_count\_anova <-   
 anova\_test(df\_long\_ext\_fix\_count,   
 fix\_count ~ iu\_group \* stimulus \* time + Error(id/(stimulus\*time)),  
 effect.size = "pes")   
  
# obtain the mixed ANOVA results  
get\_anova\_table(ext\_fix\_count\_anova)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 iu\_group 1 137 7.672 0.006 \* 0.053000  
## 2 stimulus 1 137 4.155 0.043 \* 0.029000  
## 3 time 1 137 5.733 0.018 \* 0.040000  
## 4 iu\_group:stimulus 1 137 3.460 0.065 0.025000  
## 5 iu\_group:time 1 137 4.572 0.034 \* 0.032000  
## 6 stimulus:time 1 137 0.061 0.806 0.000443  
## 7 iu\_group:stimulus:time 1 137 0.600 0.440 0.004000

# results:  
# IU: F(1,137) = 7.67, p = .006 \*\*\*, eta2(partial) = .053  
# Stimulus: F(1,137) = 4.16, p = .043 \*, eta2(partial) = .029  
# Time: F(1,137) = 5.73, p = .018 \*, eta2(partial) = .049  
# IU \* Stimulus: F(1, 137) = 3.46, p = .065, eta2(partial) = .025  
# IU \* Time: F(1,137) = 4.57, p = .034 \*, eta2(partial) = . 032  
# Stimulus \* Time: F(1,137) = 0.06, p = .806, eta2(partial) < .001  
# IU \* Stimulus \* Time: F(1,137) = 0.60, p = .440, eta2(partial) = .004  
  
# therefore, there is a significant effect of IU, Stimulus and Time on fixation count in extinction,   
# as well as a significant interaction effect of IU \* Time,  
# but no other significant interactions.   
  
# write to csv  
write.csv((get\_anova\_table(ext\_fix\_count\_anova)),   
 file = "tables/anovas/ext\_fix\_count\_anova.csv")  
  
# as there was a significant IU\*Time interaction, conduct simple   
# main effects analysis:  
## obtain effect of IU at each level of time   
simple\_effects\_ext\_fix\_count\_iu <- df\_long\_ext\_fix\_count %>%  
 group\_by(time) %>%  
 anova\_test(dv = fix\_count, wid = id, between = iu\_group, within = stimulus, effect.size = "pes") %>%  
 get\_anova\_table() %>%  
 adjust\_pvalue(method = "bonferroni")  
   
  
# get the output  
simple\_effects\_ext\_fix\_count\_iu

## # A tibble: 6 × 9  
## time Effect DFn DFd F p `p<.05` pes p.adj  
## \* <fct> <chr> <dbl> <dbl> <dbl> <dbl> <chr> <dbl> <dbl>  
## 1 -1 iu\_group 1 137 11.4 0.000952 "\*" 0.077 0.00571  
## 2 -1 stimulus 1 137 3.38 0.068 "" 0.024 0.408   
## 3 -1 iu\_group:stimulus 1 137 0.864 0.354 "" 0.006 1   
## 4 1 iu\_group 1 137 3.63 0.059 "" 0.026 0.354   
## 5 1 stimulus 1 137 1.50 0.222 "" 0.011 1   
## 6 1 iu\_group:stimulus 1 137 3.04 0.084 "" 0.022 0.504

# results:   
# the effect of IU group at early extinction was significant [F(1,137) = 11.41, p = .006, pes = .077]  
# The effect of IU group at late extinction was not significant [F(1,137) = 3.63, p = .354, pes = .026]

## *ANOVA Extinction Fixation Duration (Log Transformed)*

# transform wide format data into long format for mixed ANOVA   
df\_long\_ext\_fix\_duration\_log <- melt(df, id = c("id", "iu\_group"),   
 measure.vars = c("e\_ext\_csp\_fix\_duration\_log",   
 "e\_ext\_csm\_fix\_duration\_log",  
 "l\_ext\_csp\_fix\_duration\_log",   
 "l\_ext\_csm\_fix\_duration\_log"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_ext\_fix\_duration\_log) = c("id", "iu\_group", "condition", "fix\_duration\_log")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_ext\_fix\_duration\_log$stimulus <-   
 factor(ifelse(df\_long\_ext\_fix\_duration\_log$condition == "e\_ext\_csp\_fix\_duration\_log" |   
 df\_long\_ext\_fix\_duration\_log$condition == "l\_ext\_csp\_fix\_duration\_log", 1, -1))  
  
# create column to code extinction as early (1) and late (-1)  
df\_long\_ext\_fix\_duration\_log$time <-   
 factor(ifelse(df\_long\_ext\_fix\_duration\_log$condition == "e\_ext\_csp\_fix\_duration\_log" |   
 df\_long\_ext\_fix\_duration\_log$condition == "e\_ext\_csm\_fix\_duration\_log", 1, -1))  
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) x 2 (Time: Early, Late) mixed ANOVA,   
# and obtain effect size (partial eta squared)  
ext\_fix\_duration\_anova\_log <-   
 anova\_test(df\_long\_ext\_fix\_duration\_log,   
 fix\_duration\_log ~ iu\_group \* stimulus \* time + Error(id/(stimulus\*time)),  
 effect.size = "pes")   
  
# obtain the mixed ANOVA results  
get\_anova\_table(ext\_fix\_duration\_anova\_log)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 iu\_group 1 137 11.213 0.001 \* 0.076  
## 2 stimulus 1 137 0.510 0.477 0.004  
## 3 time 1 137 4.351 0.039 \* 0.031  
## 4 iu\_group:stimulus 1 137 5.823 0.017 \* 0.041  
## 5 iu\_group:time 1 137 0.241 0.624 0.002  
## 6 stimulus:time 1 137 0.171 0.680 0.001  
## 7 iu\_group:stimulus:time 1 137 0.946 0.333 0.007

# results:  
# IU: F(1,137) = 11.21, p < .001 \*, eta2(partial) = .076  
# Stimulus: F(1,137) = 0.51, p = .477, eta2(partial) = .004  
# Time: F(1,137) = 4.35, p = .039\*, eta2(partial) = .031  
# IU \* Stimulus: F(1, 137) = 5.82, p = .017\*, eta2(partial) = .041  
# IU \* Time: F(1,137) = 0.24, p = .624, eta2(partial) = .002  
# Stimulus \* Time: F(1,137) = 0.17, p = 680, eta2(partial) = .001  
# IU \* Stimulus \* Time: F(1,137) = 0.95, p = .333, eta2(partial) = .007  
  
# therefore, there is a significant effect of IU, Time and IU-Stimulus  
# interaction on fixation duration in extinction,   
# and no other significant effects or interactions.   
  
# write to csv  
write.csv((get\_anova\_table(ext\_fix\_duration\_anova\_log)),   
 file = "tables/anovas/ext\_fix\_duration\_anova\_log.csv")  
  
# as there was a significant IU\*Stimulus interaction, conduct simple   
# main effects analysis:  
## obtain effect of IU at each level of stimulus   
simple\_effects\_ext\_fix\_duration\_log\_iu <- df\_long\_ext\_fix\_duration\_log %>%  
 group\_by(stimulus) %>%  
 anova\_test(dv = fix\_duration\_log, wid = id, between = iu\_group, within = time, effect.size = "pes") %>%  
 get\_anova\_table() %>%  
 adjust\_pvalue(method = "bonferroni")  
  
# get the output  
simple\_effects\_ext\_fix\_duration\_log\_iu

## # A tibble: 6 × 9  
## stimulus Effect DFn DFd F p `p<.05` pes p.adj  
## \* <fct> <chr> <dbl> <dbl> <dbl> <dbl> <chr> <dbl> <dbl>  
## 1 -1 iu\_group 1 137 14.4 0.000218 "\*" 0.095 0.00131  
## 2 -1 time 1 137 4.34 0.039 "\*" 0.031 0.234   
## 3 -1 iu\_group:time 1 137 0.026 0.871 "" 0.000192 1   
## 4 1 iu\_group 1 137 6.70 0.011 "\*" 0.047 0.066   
## 5 1 time 1 137 1.94 0.166 "" 0.014 0.996   
## 6 1 iu\_group:time 1 137 0.816 0.368 "" 0.006 1

# results:   
# The effect of IU group in response to CS+ was not significant [F(1,137) = 6.70, p = .066, pes = .047]  
# the effect of IU group in response to CS- was significant [F(1,137) = 14.43, p = .001, pes = .095]

## *ANOVA Extinction Saccade Amplitude*

# transform wide format data into long format for mixed ANOVA   
df\_long\_ext\_sacc\_amplitude <- melt(df, id = c("id", "iu\_group"),   
 measure.vars = c("e\_ext\_csp\_sacc\_amplitude",   
 "e\_ext\_csm\_sacc\_amplitude",  
 "l\_ext\_csp\_sacc\_amplitude",   
 "l\_ext\_csm\_sacc\_amplitude"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_ext\_sacc\_amplitude) = c("id", "iu\_group", "condition", "sacc\_amplitude")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_ext\_sacc\_amplitude$stimulus <-   
 factor(ifelse(df\_long\_ext\_sacc\_amplitude$condition == "e\_ext\_csp\_sacc\_amplitude" |   
 df\_long\_ext\_sacc\_amplitude$condition == "l\_ext\_csp\_sacc\_amplitude", 1, -1))  
  
# create column to code extinction as early (1) and late (-1)  
df\_long\_ext\_sacc\_amplitude$time <-   
 factor(ifelse(df\_long\_ext\_sacc\_amplitude$condition == "e\_ext\_csp\_sacc\_amplitude" |   
 df\_long\_ext\_sacc\_amplitude$condition == "e\_ext\_csm\_sacc\_amplitude", 1, -1))  
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) x 2 (Time: Early, Late) mixed ANOVA,   
# and obtain effect size (partial eta squared)  
ext\_sacc\_amplitude\_anova <-   
 anova\_test(df\_long\_ext\_sacc\_amplitude,   
 sacc\_amplitude ~ iu\_group \* stimulus \* time + Error(id/(stimulus\*time)),  
 effect.size = "pes")

## Warning: NA detected in rows: 116,181,301.  
## Removing this rows before the analysis.

# obtain the mixed ANOVA results  
get\_anova\_table(ext\_sacc\_amplitude\_anova)

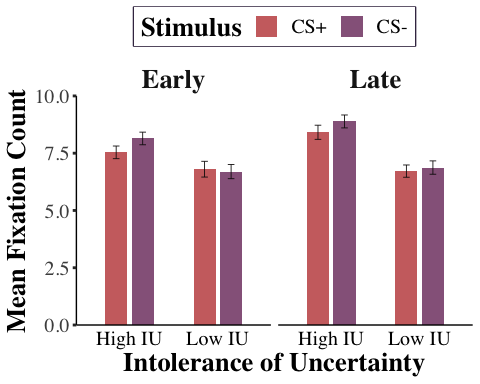
## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 iu\_group 1 134 3.170 0.077 0.023000  
## 2 stimulus 1 134 0.740 0.391 0.005000  
## 3 time 1 134 0.275 0.601 0.002000  
## 4 iu\_group:stimulus 1 134 1.687 0.196 0.012000  
## 5 iu\_group:time 1 134 0.131 0.718 0.000977  
## 6 stimulus:time 1 134 0.077 0.781 0.000577  
## 7 iu\_group:stimulus:time 1 134 0.609 0.437 0.005000

# results:  
# IU: F(1,134) = 3.17, p = .077, eta2(partial) = .023  
# Stimulus: F(1,134) = 0.74, p = .391, eta2(partial) = .005  
# Time: F(1,134) = 0.28, p = .601, eta2(partial) = .002  
# IU \* Stimulus: F(1, 134) = 1.69, p = .196, eta2(partial) = .012  
# IU \* Time: F(1,134) = 0.13, p = .718, eta2(partial) < .001  
# Stimulus \* Time: F(1,134) = 0.08, p = .781, eta2(partial) < .001  
# IU \* Stimulus \* Time: F(1,134) = .61, p = .437, eta2(partial) < .001  
  
# therefore, there are no significant effects or interactions   
# on saccade amplitude throughout extinction   
  
# write to csv  
write.csv((get\_anova\_table(ext\_sacc\_amplitude\_anova)),   
 file = "tables/anovas/ext\_sacc\_amplitude\_anova.csv")

# *Bar Graphs - Extinction Only*

## *Fixation Count*

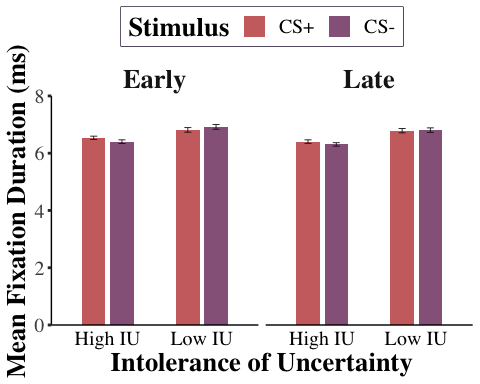
# obtain mean fix count for each group at each stimulus type and save as vector   
mean\_e\_ext\_fix\_count\_high\_iu\_csp <-   
 mean(df$e\_ext\_csp\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1"], na.rm = TRUE) # high IU CS+ early  
mean\_e\_ext\_fix\_count\_low\_iu\_csp <-   
 mean(df$e\_ext\_csp\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1"], na.rm = TRUE) # low IU CS+ early  
mean\_e\_ext\_fix\_count\_high\_iu\_csm <-   
 mean(df$e\_ext\_csm\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1"], na.rm = TRUE) # high IU CS- early  
mean\_e\_ext\_fix\_count\_low\_iu\_csm <-   
 mean(df$e\_ext\_csm\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1"], na.rm = TRUE) # low IU CS- early  
mean\_l\_ext\_fix\_count\_high\_iu\_csp <-   
 mean(df$l\_ext\_csp\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1"], na.rm = TRUE) # high IU CS+ late  
mean\_l\_ext\_fix\_count\_low\_iu\_csp <-   
 mean(df$l\_ext\_csp\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1"], na.rm = TRUE) # low IU CS+ late  
mean\_l\_ext\_fix\_count\_high\_iu\_csm <-   
 mean(df$l\_ext\_csm\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1"], na.rm = TRUE) # high IU CS- late  
mean\_l\_ext\_fix\_count\_low\_iu\_csm <-   
 mean(df$l\_ext\_csm\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1"], na.rm = TRUE) # low IU CS- late  
  
# combine into single variable   
all\_mean\_ext\_fix\_count <-   
 c(mean\_e\_ext\_fix\_count\_high\_iu\_csp, mean\_e\_ext\_fix\_count\_low\_iu\_csp,   
 mean\_e\_ext\_fix\_count\_high\_iu\_csm, mean\_e\_ext\_fix\_count\_low\_iu\_csm,  
 mean\_l\_ext\_fix\_count\_high\_iu\_csp, mean\_l\_ext\_fix\_count\_low\_iu\_csp,   
 mean\_l\_ext\_fix\_count\_high\_iu\_csm, mean\_l\_ext\_fix\_count\_low\_iu\_csm)  
  
# obtain SD fix count for each group at each stimulus type and save as vector   
sd\_e\_ext\_fix\_count\_high\_iu\_csp <-   
 sd(df$e\_ext\_csp\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1"], na.rm = TRUE) # high IU CS+ early  
sd\_e\_ext\_fix\_count\_low\_iu\_csp <-   
 sd(df$e\_ext\_csp\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1"], na.rm = TRUE) # low IU CS+ early  
sd\_e\_ext\_fix\_count\_high\_iu\_csm <-   
 sd(df$e\_ext\_csm\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1"], na.rm = TRUE) # high IU CS- early  
sd\_e\_ext\_fix\_count\_low\_iu\_csm <-   
 sd(df$e\_ext\_csm\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1"], na.rm = TRUE) # low IU CS- early  
sd\_l\_ext\_fix\_count\_high\_iu\_csp <-   
 sd(df$l\_ext\_csp\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1"], na.rm = TRUE) # high IU CS+ late  
sd\_l\_ext\_fix\_count\_low\_iu\_csp <-   
 sd(df$l\_ext\_csp\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1"], na.rm = TRUE) # low IU CS+ late  
sd\_l\_ext\_fix\_count\_high\_iu\_csm <-   
 sd(df$l\_ext\_csm\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1"], na.rm = TRUE) # high IU CS- late  
sd\_l\_ext\_fix\_count\_low\_iu\_csm <-   
 sd(df$l\_ext\_csm\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1"], na.rm = TRUE) # low IU CS- late  
  
# obtain SE:  
se\_e\_ext\_fix\_count\_high\_iu\_csp <- sd\_e\_ext\_fix\_count\_high\_iu\_csp/sqrt(length(df$id))  
se\_e\_ext\_fix\_count\_low\_iu\_csp <- sd\_e\_ext\_fix\_count\_low\_iu\_csp/sqrt(length(df$id))  
se\_e\_ext\_fix\_count\_high\_iu\_csm <- sd\_e\_ext\_fix\_count\_high\_iu\_csm/sqrt(length(df$id))  
se\_e\_ext\_fix\_count\_low\_iu\_csm <- sd\_e\_ext\_fix\_count\_low\_iu\_csm/sqrt(length(df$id))  
se\_l\_ext\_fix\_count\_high\_iu\_csp <- sd\_l\_ext\_fix\_count\_high\_iu\_csp/sqrt(length(df$id))  
se\_l\_ext\_fix\_count\_low\_iu\_csp <- sd\_l\_ext\_fix\_count\_low\_iu\_csp/sqrt(length(df$id))  
se\_l\_ext\_fix\_count\_high\_iu\_csm <- sd\_l\_ext\_fix\_count\_high\_iu\_csm/sqrt(length(df$id))  
se\_l\_ext\_fix\_count\_low\_iu\_csm <- sd\_l\_ext\_fix\_count\_low\_iu\_csm/sqrt(length(df$id))  
  
# Combine all into single variable called all\_se  
all\_se\_ext\_fix\_count <- c(se\_e\_ext\_fix\_count\_high\_iu\_csp, se\_e\_ext\_fix\_count\_low\_iu\_csp,  
 se\_e\_ext\_fix\_count\_high\_iu\_csm, se\_e\_ext\_fix\_count\_low\_iu\_csm,  
 se\_l\_ext\_fix\_count\_high\_iu\_csp, se\_l\_ext\_fix\_count\_low\_iu\_csp,   
 se\_l\_ext\_fix\_count\_high\_iu\_csm, se\_l\_ext\_fix\_count\_low\_iu\_csm)  
  
### Create new data frame for figures  
# Which includes mean and SE for each condition  
df\_fig\_extinction\_fix\_count <- data.frame(all\_mean\_ext\_fix\_count, all\_se\_ext\_fix\_count)  
  
### add labels   
# add two more variables to indicate IU group and stimulus type.   
# for IU group  
df\_fig\_extinction\_fix\_count$iu\_group[1] <- "High IU"  
df\_fig\_extinction\_fix\_count$iu\_group[2] <- "Low IU"  
df\_fig\_extinction\_fix\_count$iu\_group[3] <- "High IU"  
df\_fig\_extinction\_fix\_count$iu\_group[4] <- "Low IU"  
df\_fig\_extinction\_fix\_count$iu\_group[5] <- "High IU"  
df\_fig\_extinction\_fix\_count$iu\_group[6] <- "Low IU"  
df\_fig\_extinction\_fix\_count$iu\_group[7] <- "High IU"  
df\_fig\_extinction\_fix\_count$iu\_group[8] <- "Low IU"  
  
# for stimulus  
df\_fig\_extinction\_fix\_count$stimulus[1] <- "CS+"  
df\_fig\_extinction\_fix\_count$stimulus[2] <- "CS+"  
df\_fig\_extinction\_fix\_count$stimulus[3] <- "CS-"  
df\_fig\_extinction\_fix\_count$stimulus[4] <- "CS-"  
df\_fig\_extinction\_fix\_count$stimulus[5] <- "CS+"  
df\_fig\_extinction\_fix\_count$stimulus[6] <- "CS+"  
df\_fig\_extinction\_fix\_count$stimulus[7] <- "CS-"  
df\_fig\_extinction\_fix\_count$stimulus[8] <- "CS-"  
  
# and re-order levels of stimulus factor so that CS+ appears on left in the graph  
df\_fig\_extinction\_fix\_count$stimulus <-   
 factor(df\_fig\_extinction\_fix\_count$stimulus,levels=c("CS+","CS-"))  
  
# for early / late extinction  
df\_fig\_extinction\_fix\_count$time[1] <- "Early"  
df\_fig\_extinction\_fix\_count$time[2] <- "Early"  
df\_fig\_extinction\_fix\_count$time[3] <- "Early"  
df\_fig\_extinction\_fix\_count$time[4] <- "Early"  
df\_fig\_extinction\_fix\_count$time[5] <- "Late"  
df\_fig\_extinction\_fix\_count$time[6] <- "Late"  
df\_fig\_extinction\_fix\_count$time[7] <- "Late"  
df\_fig\_extinction\_fix\_count$time[8] <- "Late"  
  
### create figure  
fig\_extinction\_fix\_count <-   
 ggplot(df\_fig\_extinction\_fix\_count, aes(x = iu\_group, y = all\_mean\_ext\_fix\_count,   
 fill = stimulus)) +   
 geom\_bar(stat = "identity", position = position\_dodge(.6), width = .5, alpha = .85) +  
 scale\_y\_continuous(limits = c(0, 10), expand = c(0,0)) +  
 facet\_wrap(~ time) +  
 theme\_classic() +   
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_blank()) +  
 theme(axis.text.y = element\_text(size = 15), axis.ticks.y = element\_line(size = 1),   
 axis.line.y = element\_line(colour = "black")) +  
 theme(axis.text.x = element\_text(colour = "black", size = 15),   
 axis.ticks.x = element\_blank(),  
 axis.line.x = element\_line(colour = "black")) +  
 theme(axis.title = element\_text(size = 20, face = "bold")) +  
 theme(legend.position = "top",   
 legend.title = element\_text(size = 20, face = "bold"),   
 legend.box.background = element\_rect(size = .75, colour = "#403250")) +  
 theme(legend.text = element\_text(size = 15)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372")) +  
 labs(fill = "Stimulus") +  
 labs(y = "Mean Fixation Count", x = "Intolerance of Uncertainty") +  
 geom\_errorbar(aes(ymin = all\_mean\_ext\_fix\_count - all\_se\_ext\_fix\_count,   
 ymax = all\_mean\_ext\_fix\_count + all\_se\_ext\_fix\_count),   
 width = .15, position = position\_dodge(.6), colour = "#090707", size = .3) +  
 theme(strip.background = element\_blank()) +  
 theme(strip.text = element\_text(size = 20, face = "bold"))   
  
 # obtain and check figure   
print(fig\_extinction\_fix\_count)



# save figure to files  
ggsave(filename = "graphs/bar\_plots/extinction\_fix\_count.png",   
 plot = fig\_extinction\_fix\_count,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

## *Fixation Duration (Log Transformed)*

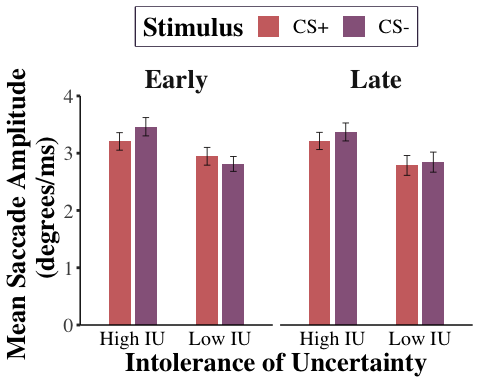
# obtain mean fix duration for each group at each stimulus type and save as vector   
# high IU CS+ early  
mean\_e\_ext\_fix\_duration\_high\_iu\_csp\_log <-   
 mean(df$e\_ext\_csp\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS+ early  
mean\_e\_ext\_fix\_duration\_low\_iu\_csp\_log <-   
 mean(df$e\_ext\_csp\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS- early  
mean\_e\_ext\_fix\_duration\_high\_iu\_csm\_log <-   
 mean(df$e\_ext\_csm\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS- early  
mean\_e\_ext\_fix\_duration\_low\_iu\_csm\_log <-   
 mean(df$e\_ext\_csm\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS+ late  
mean\_l\_ext\_fix\_duration\_high\_iu\_csp\_log <-   
 mean(df$l\_ext\_csp\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS+ late  
mean\_l\_ext\_fix\_duration\_low\_iu\_csp\_log <-   
 mean(df$l\_ext\_csp\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS- late  
mean\_l\_ext\_fix\_duration\_high\_iu\_csm\_log <-   
 mean(df$l\_ext\_csm\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS- late  
mean\_l\_ext\_fix\_duration\_low\_iu\_csm\_log <-   
 mean(df$l\_ext\_csm\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1"], na.rm = TRUE)   
  
# combine into single variable called   
all\_mean\_ext\_fix\_duration\_log <-   
 c(mean\_e\_ext\_fix\_duration\_high\_iu\_csp\_log, mean\_e\_ext\_fix\_duration\_low\_iu\_csp\_log,   
 mean\_e\_ext\_fix\_duration\_high\_iu\_csm\_log, mean\_e\_ext\_fix\_duration\_low\_iu\_csm\_log,  
 mean\_l\_ext\_fix\_duration\_high\_iu\_csp\_log, mean\_l\_ext\_fix\_duration\_low\_iu\_csp\_log,   
 mean\_l\_ext\_fix\_duration\_high\_iu\_csm\_log, mean\_l\_ext\_fix\_duration\_low\_iu\_csm\_log)  
  
# obtain SD fix duration for each group at each stimulus type and save as vector   
# high IU CS+ early  
sd\_e\_ext\_fix\_duration\_high\_iu\_csp\_log <-   
 sd(df$e\_ext\_csp\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS+ early  
sd\_e\_ext\_fix\_duration\_low\_iu\_csp\_log <-   
 sd(df$e\_ext\_csp\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS- early  
sd\_e\_ext\_fix\_duration\_high\_iu\_csm\_log <-   
 sd(df$e\_ext\_csm\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS- early  
sd\_e\_ext\_fix\_duration\_low\_iu\_csm\_log <-   
 sd(df$e\_ext\_csm\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS+ late  
sd\_l\_ext\_fix\_duration\_high\_iu\_csp\_log <-   
 sd(df$l\_ext\_csp\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS+ late  
sd\_l\_ext\_fix\_duration\_low\_iu\_csp\_log <-   
 sd(df$l\_ext\_csp\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS- late  
sd\_l\_ext\_fix\_duration\_high\_iu\_csm\_log <-   
 sd(df$l\_ext\_csm\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS- late  
sd\_l\_ext\_fix\_duration\_low\_iu\_csm\_log <-   
 sd(df$l\_ext\_csm\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1"], na.rm = TRUE)   
  
# obtain SE:  
se\_e\_ext\_fix\_duration\_high\_iu\_csp\_log <- sd\_e\_ext\_fix\_duration\_high\_iu\_csp\_log/sqrt(length(df$id))  
se\_e\_ext\_fix\_duration\_low\_iu\_csp\_log <- sd\_e\_ext\_fix\_duration\_low\_iu\_csp\_log/sqrt(length(df$id))  
se\_e\_ext\_fix\_duration\_high\_iu\_csm\_log <- sd\_e\_ext\_fix\_duration\_high\_iu\_csm\_log/sqrt(length(df$id))  
se\_e\_ext\_fix\_duration\_low\_iu\_csm\_log <- sd\_e\_ext\_fix\_duration\_low\_iu\_csm\_log/sqrt(length(df$id))  
se\_l\_ext\_fix\_duration\_high\_iu\_csp\_log <- sd\_l\_ext\_fix\_duration\_high\_iu\_csp\_log/sqrt(length(df$id))  
se\_l\_ext\_fix\_duration\_low\_iu\_csp\_log <- sd\_l\_ext\_fix\_duration\_low\_iu\_csp\_log/sqrt(length(df$id))  
se\_l\_ext\_fix\_duration\_high\_iu\_csm\_log <- sd\_l\_ext\_fix\_duration\_high\_iu\_csm\_log/sqrt(length(df$id))  
se\_l\_ext\_fix\_duration\_low\_iu\_csm\_log <- sd\_l\_ext\_fix\_duration\_low\_iu\_csm\_log/sqrt(length(df$id))  
  
# combine all into single variable   
all\_se\_ext\_fix\_duration\_log <- c(se\_e\_ext\_fix\_duration\_high\_iu\_csp\_log, se\_e\_ext\_fix\_duration\_low\_iu\_csp\_log,  
 se\_e\_ext\_fix\_duration\_high\_iu\_csm\_log, se\_e\_ext\_fix\_duration\_low\_iu\_csm\_log,  
 se\_l\_ext\_fix\_duration\_high\_iu\_csp\_log, se\_l\_ext\_fix\_duration\_low\_iu\_csp\_log,   
 se\_l\_ext\_fix\_duration\_high\_iu\_csm\_log, se\_l\_ext\_fix\_duration\_low\_iu\_csm\_log)  
  
# create new data frame for figures which includes mean and SE for each condition  
df\_fig\_extinction\_fix\_duration\_log <- data.frame(all\_mean\_ext\_fix\_duration\_log, all\_se\_ext\_fix\_duration\_log)  
  
# add labels - add two more variables to indicate IU group, stimulus type and extinction time   
# for IU group  
df\_fig\_extinction\_fix\_duration\_log$iu\_group[1] <- "High IU"  
df\_fig\_extinction\_fix\_duration\_log$iu\_group[2] <- "Low IU"  
df\_fig\_extinction\_fix\_duration\_log$iu\_group[3] <- "High IU"  
df\_fig\_extinction\_fix\_duration\_log$iu\_group[4] <- "Low IU"  
df\_fig\_extinction\_fix\_duration\_log$iu\_group[5] <- "High IU"  
df\_fig\_extinction\_fix\_duration\_log$iu\_group[6] <- "Low IU"  
df\_fig\_extinction\_fix\_duration\_log$iu\_group[7] <- "High IU"  
df\_fig\_extinction\_fix\_duration\_log$iu\_group[8] <- "Low IU"  
  
# for stimulus  
df\_fig\_extinction\_fix\_duration\_log$stimulus[1] <- "CS+"  
df\_fig\_extinction\_fix\_duration\_log$stimulus[2] <- "CS+"  
df\_fig\_extinction\_fix\_duration\_log$stimulus[3] <- "CS-"  
df\_fig\_extinction\_fix\_duration\_log$stimulus[4] <- "CS-"  
df\_fig\_extinction\_fix\_duration\_log$stimulus[5] <- "CS+"  
df\_fig\_extinction\_fix\_duration\_log$stimulus[6] <- "CS+"  
df\_fig\_extinction\_fix\_duration\_log$stimulus[7] <- "CS-"  
df\_fig\_extinction\_fix\_duration\_log$stimulus[8] <- "CS-"  
  
# and re-order levels of stimulus factor so that CS+ appears on left in the graph  
df\_fig\_extinction\_fix\_duration\_log$stimulus <-   
 factor(df\_fig\_extinction\_fix\_duration\_log$stimulus,levels=c("CS+","CS-"))  
  
# for early / late extinction  
df\_fig\_extinction\_fix\_duration\_log$time[1] <- "Early"  
df\_fig\_extinction\_fix\_duration\_log$time[2] <- "Early"  
df\_fig\_extinction\_fix\_duration\_log$time[3] <- "Early"  
df\_fig\_extinction\_fix\_duration\_log$time[4] <- "Early"  
df\_fig\_extinction\_fix\_duration\_log$time[5] <- "Late"  
df\_fig\_extinction\_fix\_duration\_log$time[6] <- "Late"  
df\_fig\_extinction\_fix\_duration\_log$time[7] <- "Late"  
df\_fig\_extinction\_fix\_duration\_log$time[8] <- "Late"  
  
# create figure  
fig\_extinction\_fix\_duration\_log <-   
 ggplot(df\_fig\_extinction\_fix\_duration\_log, aes(x = iu\_group, y = all\_mean\_ext\_fix\_duration\_log,   
 fill = stimulus)) +   
 geom\_bar(stat = "identity", position = position\_dodge(.6), width = .5, alpha = .85) +  
 scale\_y\_continuous(limits = c(0, 8), expand = c(0,0)) +  
 facet\_wrap(~ time) +  
 theme\_classic() +   
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_blank()) +  
 theme(axis.text.y = element\_text(size = 15), axis.ticks.y = element\_line(size = 1),   
 axis.line.y = element\_line(colour = "black")) +  
 theme(axis.text.x = element\_text(colour = "black", size = 15),   
 axis.ticks.x = element\_blank(),  
 axis.line.x = element\_line(colour = "black")) +  
 theme(axis.title = element\_text(size = 20, face = "bold")) +  
 theme(legend.position = "top",   
 legend.title = element\_text(size = 20, face = "bold"),   
 legend.box.background = element\_rect(size = .75, colour = "#403250")) +  
 theme(legend.text = element\_text(size = 15)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372")) +  
 labs(fill = "Stimulus") +  
 labs(y = "Mean Fixation Duration (ms)", x = "Intolerance of Uncertainty") +  
 geom\_errorbar(aes(ymin = all\_mean\_ext\_fix\_duration\_log - all\_se\_ext\_fix\_duration\_log,   
 ymax = all\_mean\_ext\_fix\_duration\_log + all\_se\_ext\_fix\_duration\_log),   
 width = .15, position = position\_dodge(.6), colour = "#090707", size = .3) +  
 theme(strip.background = element\_blank()) +  
 theme(strip.text = element\_text(size = 20, face = "bold"))   
  
  
# obtain and check figure   
print(fig\_extinction\_fix\_duration\_log)



# save figure to files  
ggsave(filename = "graphs/bar\_plots/extinction\_fix\_duration\_log.png",   
 plot = fig\_extinction\_fix\_duration\_log,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

## *Saccade Amplitude*

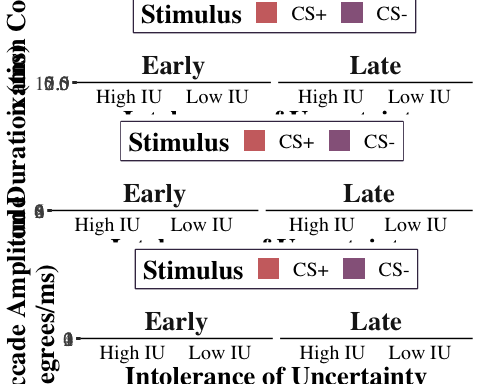
# obtain mean sacc amplitude for each group at each stimulus type and save as vector   
# high IU CS+ early  
mean\_e\_ext\_sacc\_amplitude\_high\_iu\_csp <-   
 mean(df$e\_ext\_csp\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS+ early  
mean\_e\_ext\_sacc\_amplitude\_low\_iu\_csp <-   
 mean(df$e\_ext\_csp\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS- early  
mean\_e\_ext\_sacc\_amplitude\_high\_iu\_csm <-   
 mean(df$e\_ext\_csm\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS- early  
mean\_e\_ext\_sacc\_amplitude\_low\_iu\_csm <-   
 mean(df$e\_ext\_csm\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS+ late  
mean\_l\_ext\_sacc\_amplitude\_high\_iu\_csp <-   
 mean(df$l\_ext\_csp\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS+ late  
mean\_l\_ext\_sacc\_amplitude\_low\_iu\_csp <-   
 mean(df$l\_ext\_csp\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS- late  
mean\_l\_ext\_sacc\_amplitude\_high\_iu\_csm <-   
 mean(df$l\_ext\_csm\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS- late  
mean\_l\_ext\_sacc\_amplitude\_low\_iu\_csm <-   
 mean(df$l\_ext\_csm\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1"], na.rm = TRUE)   
  
# combine into single variable called   
all\_mean\_ext\_sacc\_amplitude <-   
 c(mean\_e\_ext\_sacc\_amplitude\_high\_iu\_csp, mean\_e\_ext\_sacc\_amplitude\_low\_iu\_csp,   
 mean\_e\_ext\_sacc\_amplitude\_high\_iu\_csm, mean\_e\_ext\_sacc\_amplitude\_low\_iu\_csm,  
 mean\_l\_ext\_sacc\_amplitude\_high\_iu\_csp, mean\_l\_ext\_sacc\_amplitude\_low\_iu\_csp,   
 mean\_l\_ext\_sacc\_amplitude\_high\_iu\_csm, mean\_l\_ext\_sacc\_amplitude\_low\_iu\_csm)  
  
# obtain SD sacc amplitude for each group at each stimulus type and save as vector   
# high IU CS+ early  
sd\_e\_ext\_sacc\_amplitude\_high\_iu\_csp <-   
 sd(df$e\_ext\_csp\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS+ early  
sd\_e\_ext\_sacc\_amplitude\_low\_iu\_csp <-   
 sd(df$e\_ext\_csp\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS- early  
sd\_e\_ext\_sacc\_amplitude\_high\_iu\_csm <-   
 sd(df$e\_ext\_csm\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS- early  
sd\_e\_ext\_sacc\_amplitude\_low\_iu\_csm <-   
 sd(df$e\_ext\_csm\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS+ late  
sd\_l\_ext\_sacc\_amplitude\_high\_iu\_csp <-   
 sd(df$l\_ext\_csp\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS+ late  
sd\_l\_ext\_sacc\_amplitude\_low\_iu\_csp <-   
 sd(df$l\_ext\_csp\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1"], na.rm = TRUE)   
  
# high IU CS- late  
sd\_l\_ext\_sacc\_amplitude\_high\_iu\_csm <-   
 sd(df$l\_ext\_csm\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1"], na.rm = TRUE)   
  
# low IU CS- late  
sd\_l\_ext\_sacc\_amplitude\_low\_iu\_csm <-   
 sd(df$l\_ext\_csm\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1"], na.rm = TRUE)   
  
# obtain SE:  
se\_e\_ext\_sacc\_amplitude\_high\_iu\_csp <- sd\_e\_ext\_sacc\_amplitude\_high\_iu\_csp/sqrt(length(df$id))  
se\_e\_ext\_sacc\_amplitude\_low\_iu\_csp <- sd\_e\_ext\_sacc\_amplitude\_low\_iu\_csp/sqrt(length(df$id))  
se\_e\_ext\_sacc\_amplitude\_high\_iu\_csm <- sd\_e\_ext\_sacc\_amplitude\_high\_iu\_csm/sqrt(length(df$id))  
se\_e\_ext\_sacc\_amplitude\_low\_iu\_csm <- sd\_e\_ext\_sacc\_amplitude\_low\_iu\_csm/sqrt(length(df$id))  
se\_l\_ext\_sacc\_amplitude\_high\_iu\_csp <- sd\_l\_ext\_sacc\_amplitude\_high\_iu\_csp/sqrt(length(df$id))  
se\_l\_ext\_sacc\_amplitude\_low\_iu\_csp <- sd\_l\_ext\_sacc\_amplitude\_low\_iu\_csp/sqrt(length(df$id))  
se\_l\_ext\_sacc\_amplitude\_high\_iu\_csm <- sd\_l\_ext\_sacc\_amplitude\_high\_iu\_csm/sqrt(length(df$id))  
se\_l\_ext\_sacc\_amplitude\_low\_iu\_csm <- sd\_l\_ext\_sacc\_amplitude\_low\_iu\_csm/sqrt(length(df$id))  
  
# combine all into single variable   
all\_se\_ext\_sacc\_amplitude <- c(se\_e\_ext\_sacc\_amplitude\_high\_iu\_csp, se\_e\_ext\_sacc\_amplitude\_low\_iu\_csp,  
 se\_e\_ext\_sacc\_amplitude\_high\_iu\_csm, se\_e\_ext\_sacc\_amplitude\_low\_iu\_csm,  
 se\_l\_ext\_sacc\_amplitude\_high\_iu\_csp, se\_l\_ext\_sacc\_amplitude\_low\_iu\_csp,   
 se\_l\_ext\_sacc\_amplitude\_high\_iu\_csm, se\_l\_ext\_sacc\_amplitude\_low\_iu\_csm)  
  
# create new data frame for figures which includes mean and SE for each condition  
df\_fig\_extinction\_sacc\_amplitude <- data.frame(all\_mean\_ext\_sacc\_amplitude, all\_se\_ext\_sacc\_amplitude)  
  
# add labels - add two more variables to indicate IU group, stimulus type and extinction time   
# for IU group  
df\_fig\_extinction\_sacc\_amplitude$iu\_group[1] <- "High IU"  
df\_fig\_extinction\_sacc\_amplitude$iu\_group[2] <- "Low IU"  
df\_fig\_extinction\_sacc\_amplitude$iu\_group[3] <- "High IU"  
df\_fig\_extinction\_sacc\_amplitude$iu\_group[4] <- "Low IU"  
df\_fig\_extinction\_sacc\_amplitude$iu\_group[5] <- "High IU"  
df\_fig\_extinction\_sacc\_amplitude$iu\_group[6] <- "Low IU"  
df\_fig\_extinction\_sacc\_amplitude$iu\_group[7] <- "High IU"  
df\_fig\_extinction\_sacc\_amplitude$iu\_group[8] <- "Low IU"  
  
# for stimulus  
df\_fig\_extinction\_sacc\_amplitude$stimulus[1] <- "CS+"  
df\_fig\_extinction\_sacc\_amplitude$stimulus[2] <- "CS+"  
df\_fig\_extinction\_sacc\_amplitude$stimulus[3] <- "CS-"  
df\_fig\_extinction\_sacc\_amplitude$stimulus[4] <- "CS-"  
df\_fig\_extinction\_sacc\_amplitude$stimulus[5] <- "CS+"  
df\_fig\_extinction\_sacc\_amplitude$stimulus[6] <- "CS+"  
df\_fig\_extinction\_sacc\_amplitude$stimulus[7] <- "CS-"  
df\_fig\_extinction\_sacc\_amplitude$stimulus[8] <- "CS-"  
  
# and re-order levels of stimulus factor so that CS+ appears on left in the graph  
df\_fig\_extinction\_sacc\_amplitude$stimulus <-   
 factor(df\_fig\_extinction\_sacc\_amplitude$stimulus,levels=c("CS+","CS-"))  
  
# for early / late extinction  
df\_fig\_extinction\_sacc\_amplitude$time[1] <- "Early"  
df\_fig\_extinction\_sacc\_amplitude$time[2] <- "Early"  
df\_fig\_extinction\_sacc\_amplitude$time[3] <- "Early"  
df\_fig\_extinction\_sacc\_amplitude$time[4] <- "Early"  
df\_fig\_extinction\_sacc\_amplitude$time[5] <- "Late"  
df\_fig\_extinction\_sacc\_amplitude$time[6] <- "Late"  
df\_fig\_extinction\_sacc\_amplitude$time[7] <- "Late"  
df\_fig\_extinction\_sacc\_amplitude$time[8] <- "Late"  
  
# create figure  
fig\_extinction\_sacc\_amplitude <-   
 ggplot(df\_fig\_extinction\_sacc\_amplitude, aes(x = iu\_group, y = all\_mean\_ext\_sacc\_amplitude,   
 fill = stimulus)) +   
 geom\_bar(stat = "identity", position = position\_dodge(.6), width = .5, alpha = .85) +  
 scale\_y\_continuous(limits = c(0, 4), expand = c(0,0)) +  
 facet\_wrap(~ time) +  
 theme\_classic() +   
 theme(text = element\_text(family = "serif"),   
 plot.title = element\_blank()) +  
 theme(axis.text.y = element\_text(size = 15), axis.ticks.y = element\_line(size = 1),   
 axis.line.y = element\_line(colour = "black")) +  
 theme(axis.text.x = element\_text(colour = "black", size = 15),   
 axis.ticks.x = element\_blank(),  
 axis.line.x = element\_line(colour = "black")) +  
 theme(axis.title = element\_text(size = 20, face = "bold")) +  
 theme(legend.position = "top",   
 legend.title = element\_text(size = 20, face = "bold"),   
 legend.box.background = element\_rect(size = .75, colour = "#403250")) +  
 theme(legend.text = element\_text(size = 15)) +  
 scale\_fill\_manual(values = c("#c45150", "#824372")) +  
 labs(fill = "Stimulus") +  
 labs(y = "Mean Saccade Amplitude \n (degrees/ms)", x = "Intolerance of Uncertainty") +  
 geom\_errorbar(aes(ymin = all\_mean\_ext\_sacc\_amplitude - all\_se\_ext\_sacc\_amplitude,  
 ymax = all\_mean\_ext\_sacc\_amplitude + all\_se\_ext\_sacc\_amplitude),   
 width = .15, position = position\_dodge(.6), colour = "#090707", size = .3) +  
 theme(strip.background = element\_blank()) +  
 theme(strip.text = element\_text(size = 20, face = "bold"))   
  
# obtain and check figure   
print(fig\_extinction\_sacc\_amplitude)



# save figure to files  
ggsave(filename = "graphs/bar\_plots/extinction\_sacc\_amplitude.png",   
 plot = fig\_extinction\_sacc\_amplitude,   
 width = 20,   
 height = 10,   
 dpi = 300,   
 units = "cm")

## *Combine Bar Graphs*

all\_bar\_plots <- grid.arrange(fig\_extinction\_fix\_count,  
 fig\_extinction\_fix\_duration\_log,  
 fig\_extinction\_sacc\_amplitude,   
 ncol = 1)



# save figure to files  
ggsave(filename = "graphs/bar\_plots/all\_bar\_plots.png",   
 plot = all\_bar\_plots,   
 width = 20,   
 height = 30,   
 dpi = 300,   
 units = "cm")

# *ANCOVAs to test Specificity of IU over Trait Anxiety*

## *ANCOVA Acquisition Fixation Count*

# transform wide format data into long format for mixed ANCOVA   
df\_long\_acq\_fix\_count <- melt(df, id = c("id", "iu\_group", "sticsa\_total"),   
 measure.vars = c("acq\_csp\_fix\_count",   
 "acq\_csm\_fix\_count"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_acq\_fix\_count) = c("id", "iu\_group", "sticsa\_total", "condition", "fix\_count")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_acq\_fix\_count$stimulus <-   
 factor(ifelse(df\_long\_acq\_fix\_count$condition == "acq\_csp\_fix\_count", 1, -1))  
  
# mean centre continuous covariate (STICSA)  
# to apply mean centring, first obtain average sticsa scores for all participants,   
# and save as a variable  
df\_long\_acq\_fix\_count$sticsa\_total\_avg <- mean(df\_long\_acq\_fix\_count$sticsa\_total)  
  
# next, subtract this average from all participants' sticsa scores,  
# and save as a variable  
df\_long\_acq\_fix\_count$sticsa\_total\_centred <-   
 df\_long\_acq\_fix\_count$sticsa\_total - df\_long\_acq\_fix\_count$sticsa\_total\_avg   
# from this we have mean sticsa scores after centring   
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) mixed ANCOVA,   
# with mean-cenred STICSA as covariate  
# and obtain effect size (partial eta squared)  
acq\_fix\_count\_ancova <-   
 anova\_test(df\_long\_acq\_fix\_count, fix\_count ~ iu\_group \* stimulus + Error(id/stimulus),  
 covariate = sticsa\_total\_centred, effect.size = "pes")   
  
# obtain the mixed ANCOVA results  
get\_anova\_table(acq\_fix\_count\_ancova)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 sticsa\_total\_centred 1 136 0.059 0.808000 0.000434  
## 2 iu\_group 1 136 3.191 0.076000 0.023000  
## 3 stimulus 1 136 11.622 0.000858 \* 0.079000  
## 4 sticsa\_total\_centred:stimulus 1 136 1.845 0.177000 0.013000  
## 5 iu\_group:stimulus 1 136 1.230 0.269000 0.009000

# results:  
# STICSA (centred): F(1,136) = 0.06, p = .808, eta2(partial) = < .001  
# IU: F(1,136) = 3.19, p = .076, eta2(partial) = .023  
# Stimulus: F(1,136) = 11.62, p < .001\*\*\*, eta2(partial) = .079  
# STICSA \* Stimulus: F(1,136) = 1.85, p = .177, eta2(partial) = .013  
# IU \* Stimulus: F(1, 136) = 1.23, p = .269, eta2(partial) = .009  
  
# therefore, after accounting for trait anxiety, IU no longer has a significant  
# effect on fixation count in acquisition, but stimulus continues to have  
# significant effect. IU\*Stimulus interaction also remains non-significant,   
# even after controlling for trait anxiety.  
  
# write to csv  
write.csv((get\_anova\_table(acq\_fix\_count\_ancova)),   
 file = "tables/ancovas/acq\_fix\_count\_ancova.csv")

## *ANCOVA Acquisition Fixation Duration (Log Transformed)*

# transform wide format data into long format for mixed ANCOVA   
df\_long\_acq\_fix\_duration\_log <- melt(df, id = c("id", "iu\_group", "sticsa\_total"),   
 measure.vars = c("acq\_csp\_fix\_duration\_log",   
 "acq\_csm\_fix\_duration\_log"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_acq\_fix\_duration\_log) = c("id", "iu\_group", "sticsa\_total", "condition", "fix\_duration\_log")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_acq\_fix\_duration\_log$stimulus <-   
 factor(ifelse(df\_long\_acq\_fix\_duration\_log$condition == "acq\_csp\_fix\_duration\_log", 1, -1))  
  
# mean centre continuous covariate (STICSA)  
# to apply mean centring, first obtain average sticsa scores for all participants,   
# and save as a variable  
df\_long\_acq\_fix\_duration\_log$sticsa\_total\_avg <- mean(df\_long\_acq\_fix\_duration\_log$sticsa\_total)  
  
# next, subtract this average from all participants' sticsa scores,  
# and save as a variable  
df\_long\_acq\_fix\_duration\_log$sticsa\_total\_centred <-   
 df\_long\_acq\_fix\_duration\_log$sticsa\_total - df\_long\_acq\_fix\_duration\_log$sticsa\_total\_avg   
# from this we have mean sticsa scores after centring   
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) mixed ANCOVA,   
# with mean-cenred STICSA as covariate  
# and obtain effect size (partial eta squared)  
acq\_fix\_duration\_ancova\_log <-   
 anova\_test(df\_long\_acq\_fix\_duration\_log, fix\_duration\_log ~ iu\_group \* stimulus + Error(id/stimulus),  
 covariate = sticsa\_total\_centred, effect.size = "pes")   
  
# obtain the mixed ANCOVA results  
get\_anova\_table(acq\_fix\_duration\_ancova\_log)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 sticsa\_total\_centred 1 136 0.268 0.606 0.002  
## 2 iu\_group 1 136 3.890 0.051 0.028  
## 3 stimulus 1 136 2.935 0.089 0.021  
## 4 sticsa\_total\_centred:stimulus 1 136 0.409 0.524 0.003  
## 5 iu\_group:stimulus 1 136 1.674 0.198 0.012

# results:  
# STICSA (centred): F(1,136) = 0.27, p = .606, eta2(partial) = .002  
# IU: F(1,136) = 3.89, p = .051, eta2(partial) = .028  
# Stimulus: F(1,136) = 2.94, p = .089, eta2(partial) = .021  
# STICSA \* Stimulus: F(1,136) = 0.41, p = .524, eta2(partial) = .003  
# IU \* Stimulus: F(1, 136) = 1.67, p = .198, eta2(partial) = .012  
  
# there are no significant effects or interactions on fixation duration in acquisition.   
  
# write to csv  
write.csv((get\_anova\_table(acq\_fix\_duration\_ancova\_log)),   
 file = "tables/ancovas/acq\_fix\_duration\_ancova\_log.csv")

## *ANCOVA Acquisition Saccade Amplitude*

# transform wide format data into long format for mixed ANCOVA   
df\_long\_acq\_sacc\_amplitude <- melt(df, id = c("id", "iu\_group", "sticsa\_total"),   
 measure.vars = c("acq\_csp\_sacc\_amplitude",   
 "acq\_csm\_sacc\_amplitude"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_acq\_sacc\_amplitude) = c("id", "iu\_group", "sticsa\_total", "condition", "sacc\_amplitude")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_acq\_sacc\_amplitude$stimulus <-   
 factor(ifelse(df\_long\_acq\_sacc\_amplitude$condition == "acq\_csp\_sacc\_amplitude", 1, -1))  
  
# mean centre continuous covariate (STICSA)  
# to apply mean centring, first obtain average sticsa scores for all participants,   
# and save as a variable  
df\_long\_acq\_sacc\_amplitude$sticsa\_total\_avg <- mean(df\_long\_acq\_sacc\_amplitude$sticsa\_total)  
  
# next, subtract this average from all participants' sticsa scores,  
# and save as a variable  
df\_long\_acq\_sacc\_amplitude$sticsa\_total\_centred <-   
 df\_long\_acq\_sacc\_amplitude$sticsa\_total - df\_long\_acq\_sacc\_amplitude$sticsa\_total\_avg   
# from this we have mean sticsa scores after centring   
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) mixed ANCOVA,   
# with mean-cenred STICSA as covariate  
# and obtain effect size (partial eta squared)  
acq\_sacc\_amplitude\_ancova <-   
 anova\_test(df\_long\_acq\_sacc\_amplitude, sacc\_amplitude ~ iu\_group \* stimulus + Error(id/stimulus),  
 covariate = sticsa\_total\_centred, effect.size = "pes")

## Warning: NA detected in rows: 234,259.  
## Removing this rows before the analysis.

# obtain the mixed ANCOVA results  
get\_anova\_table(acq\_sacc\_amplitude\_ancova)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 sticsa\_total\_centred 1 134 0.007 0.935 0.0000503  
## 2 iu\_group 1 134 2.128 0.147 0.0160000  
## 3 stimulus 1 134 0.943 0.333 0.0070000  
## 4 sticsa\_total\_centred:stimulus 1 134 0.643 0.424 0.0050000  
## 5 iu\_group:stimulus 1 134 0.864 0.354 0.0060000

# results:  
# STICSA (centred): F(1,134) = 0.01, p = .935, eta2(partial) < .001  
# IU: F(1,134) = 2.13, p = .147, eta2(partial) = .016  
# Stimulus: F(1,134) = 0.94, p = .333, eta2(partial) = .007  
# STICSA \* Stimulus: F(1,134) = 0.64, p = .424, eta2(partial) = .005  
# IU \* Stimulus: F(1, 134) = 0.86, p = .354, eta2(partial) = .006  
  
# therefore, after accounting for trait anxiety, there continue not  
# to be any significant effects of IU, stimulus, and interaction  
# effects on saccade amplitude in acquisition.   
  
# write to csv  
write.csv((get\_anova\_table(acq\_sacc\_amplitude\_ancova)),   
 file = "tables/ancovas/acq\_sacc\_amplitude\_ancova.csv")

## *ANCOVA Extinction Fixation Count*

# transform wide format data into long format for mixed ANOVA   
df\_long\_ext\_fix\_count <- melt(df, id = c("id", "iu\_group", "sticsa\_total"),   
 measure.vars = c("e\_ext\_csp\_fix\_count",   
 "e\_ext\_csm\_fix\_count",  
 "l\_ext\_csp\_fix\_count",   
 "l\_ext\_csm\_fix\_count"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_ext\_fix\_count) = c("id", "iu\_group", "sticsa\_total", "condition", "fix\_count")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_ext\_fix\_count$stimulus <-   
 factor(ifelse(df\_long\_ext\_fix\_count$condition == "e\_ext\_csp\_fix\_count" |   
 df\_long\_ext\_fix\_count$condition == "l\_ext\_csp\_fix\_count", 1, -1))  
  
# create column to code extinction as early (1) and late (-1)  
df\_long\_ext\_fix\_count$time <-   
 factor(ifelse(df\_long\_ext\_fix\_count$condition == "e\_ext\_csp\_fix\_count" |   
 df\_long\_ext\_fix\_count$condition == "e\_ext\_csm\_fix\_count", 1, -1))  
  
# mean centre continuous covariate (STICSA)  
# to apply mean centring, first obtain average sticsa scores for all participants,   
# and save as a variable  
df\_long\_ext\_fix\_count$sticsa\_total\_avg <- mean(df\_long\_ext\_fix\_count$sticsa\_total)  
  
# next, subtract this average from all participants' sticsa scores,  
# and save as a variable  
df\_long\_ext\_fix\_count$sticsa\_total\_centred <-   
 df\_long\_ext\_fix\_count$sticsa\_total - df\_long\_ext\_fix\_count$sticsa\_total\_avg   
# from this we have mean sticsa scores after centring   
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) x 2 (Time: Early, Late) mixed ANOVA,   
# with mean-centred STICSA as covariate,  
# and obtain effect size (partial eta squared)  
ext\_fix\_count\_ancova <-   
 anova\_test(df\_long\_ext\_fix\_count,   
 fix\_count ~ iu\_group \* stimulus \* time + Error(id/(stimulus\*time)),  
 covariate = sticsa\_total\_centred, effect.size = "pes")   
  
# obtain the mixed ANCOVA results  
get\_anova\_table(ext\_fix\_count\_ancova)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 sticsa\_total\_centred 1 136 0.433000 0.512 0.00300000  
## 2 iu\_group 1 136 4.361000 0.039 \* 0.03100000  
## 3 stimulus 1 136 4.209000 0.042 \* 0.03000000  
## 4 time 1 136 5.692000 0.018 \* 0.04000000  
## 5 sticsa\_total\_centred:stimulus 1 136 1.098000 0.297 0.00800000  
## 6 iu\_group:stimulus 1 136 4.560000 0.035 \* 0.03200000  
## 7 sticsa\_total\_centred:time 1 136 0.000429 0.984 0.00000316  
## 8 iu\_group:time 1 136 3.489000 0.064 0.02500000  
## 9 stimulus:time 1 136 0.066000 0.797 0.00048800  
## 10 sticsa\_total\_centred:stimulus:time 1 136 0.901000 0.344 0.00700000  
## 11 iu\_group:stimulus:time 1 136 0.044000 0.834 0.00032500

# results:  
# STICSA (centred): F(1,136) = 0.43, p = .512, eta2(partial) = .003  
# IU: F(1,136) = 4.36, p = .039\*, eta2(partial) = .031  
# Stimulus: F(1,136) = 4.21, p = .042\*, eta2(partial) = .030  
# Time: F(1,136) = 5.69, p = .018 \*, eta2(partial) = .040  
# STICSA \* Stimulus: F(1,136) = 1.10, p = .297, eta2(partial) = .008  
# IU \* Stimulus: F(1, 136) = 4.56, p = .035\*, eta2(partial) = .032  
# STICSA\* Time: F(1,136) = 0.00, p = .982, eta2(partial) < .001  
# IU \* Time: F(1,136) = 3.49, p = .064, eta2(partial) = . 025  
# Stimulus \* Time: F(1,136) = 0.07, p = .797, eta2(partial) < .001  
# STICSA \* Stimulus \* Time: F(1,136) = 0.90, p = .344, eta2(partial) = .007  
# IU \* Stimulus \* Time: F(1,136) = 0.04, p = .834, eta2(partial) < .001  
  
# therefore, after accounting for trait anxiety, IU, Stimulus, and Time   
# continue to have a significant effect on fixation duration in acquisition.   
# there is no longer a significant interaction effect of IU\*Time,   
# but there is now a significant interaction effect of IU\*stimulus  
  
# write to csv  
write.csv((get\_anova\_table(ext\_fix\_count\_ancova)),   
 file = "tables/ancovas/ext\_fix\_count\_ancova.csv")  
  
# as there was a significant IU\*Stimulus interaction (which differed from observed  
# mixed ANOVA), conduct simple main effects analysis:  
## obtain effect of IU at each level of stimulus   
simple\_effects\_ext\_fix\_count\_iu\_ancova <- df\_long\_ext\_fix\_count %>%  
 group\_by(stimulus) %>%  
 anova\_test(dv = fix\_count, wid = id, between = iu\_group, within = time,  
 covariate = sticsa\_total\_centred, effect.size = "pes") %>%  
 get\_anova\_table() %>%  
 adjust\_pvalue(method = "bonferroni")  
  
# get the output  
simple\_effects\_ext\_fix\_count\_iu\_ancova

## # A tibble: 10 × 9  
## stimulus Effect DFn DFd F p `p<.05` pes p.adj  
## \* <fct> <chr> <dbl> <dbl> <dbl> <dbl> <chr> <dbl> <dbl>  
## 1 -1 sticsa\_total\_centred 1 136 0.142 0.707 "" 0.001 1   
## 2 -1 iu\_group 1 136 6.66 0.011 "\*" 0.047 0.11  
## 3 -1 time 1 136 5.02 0.027 "\*" 0.036 0.27  
## 4 -1 sticsa\_total\_centred:ti… 1 136 0.369 0.545 "" 0.003 1   
## 5 -1 iu\_group:time 1 136 2.25 0.136 "" 0.016 1   
## 6 1 sticsa\_total\_centred 1 136 0.796 0.374 "" 0.006 1   
## 7 1 iu\_group 1 136 2.16 0.143 "" 0.016 1   
## 8 1 time 1 136 2.86 0.093 "" 0.021 0.93  
## 9 1 sticsa\_total\_centred:ti… 1 136 0.253 0.616 "" 0.002 1   
## 10 1 iu\_group:time 1 136 2.40 0.124 "" 0.017 1

# results:   
# The effect of IU group on CS+ was not significant [F(1,136) = 2.17, p = .1.00, pes = .016]  
# the effect of IU group on CS- was not significant [F(1,136) = 6.66, p = .110, pes = .047]  
  
# as there was significant IU-stimulus interaction that was  
# not observed before in mixed ANOVA, obtain estimated  
# marginal means to be reported:  
  
## IU-Stimulus interaction  
# obtain emmeans  
emmeans\_ext\_fix\_count\_ancova\_iu\_stimulus <- df\_long\_ext\_fix\_count %>%  
 group\_by(stimulus) %>%  
 emmeans\_test(fix\_count ~ iu\_group, covariate = sticsa\_total\_centred) %>%  
 get\_emmeans()

## Warning: Expected 2 pieces. Additional pieces discarded in 2 rows [1, 2].

emmeans\_ext\_fix\_count\_ancova\_iu\_stimulus

## # A tibble: 4 × 9  
## sticsa\_total\_centred stimulus iu\_group emmean se df conf.low conf.high  
## <dbl> <fct> <fct> <dbl> <dbl> <dbl> <dbl> <dbl>  
## 1 -2.86e-15 -1 -1 6.88 0.303 551 6.29 7.48  
## 2 -2.86e-15 -1 1 8.41 0.311 551 7.80 9.02  
## 3 -2.86e-15 1 -1 6.86 0.303 551 6.26 7.45  
## 4 -2.86e-15 1 1 7.88 0.311 551 7.27 8.49  
## # … with 1 more variable: method <chr>

# save them as variables  
emmeans\_ext\_fix\_count\_ancova\_high\_iu\_csp <- 7.88  
emmeans\_ext\_fix\_count\_ancova\_high\_iu\_csm <- 8.41  
emmeans\_ext\_fix\_count\_ancova\_low\_iu\_csp <- 6.86  
emmeans\_ext\_fix\_count\_ancova\_low\_iu\_csm <- 6.88

## *ANCOVA Extinction Fixation Duration (Log Transformed)*

# transform wide format data into long format for mixed ANCOVA   
df\_long\_ext\_fix\_duration\_log <- melt(df, id = c("id", "iu\_group", "sticsa\_total"),   
 measure.vars = c("e\_ext\_csp\_fix\_duration\_log",   
 "e\_ext\_csm\_fix\_duration\_log",  
 "l\_ext\_csp\_fix\_duration\_log",   
 "l\_ext\_csm\_fix\_duration\_log"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_ext\_fix\_duration\_log) = c("id", "iu\_group", "sticsa\_total", "condition", "fix\_duration\_log")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_ext\_fix\_duration\_log$stimulus <-   
 factor(ifelse(df\_long\_ext\_fix\_duration\_log$condition == "e\_ext\_csp\_fix\_duration\_log" |   
 df\_long\_ext\_fix\_duration\_log$condition == "l\_ext\_csp\_fix\_duration\_log", 1, -1))  
  
# create column to code extinction as early (1) and late (-1)  
df\_long\_ext\_fix\_duration\_log$time <-   
 factor(ifelse(df\_long\_ext\_fix\_duration\_log$condition == "e\_ext\_csp\_fix\_duration\_log" |   
 df\_long\_ext\_fix\_duration\_log$condition == "e\_ext\_csm\_fix\_duration\_log", 1, -1))  
  
# mean centre continuous covariate (STICSA)  
# to apply mean centring, first obtain average sticsa scores for all participants,   
# and save as a variable  
df\_long\_ext\_fix\_duration\_log$sticsa\_total\_avg <- mean(df\_long\_ext\_fix\_duration\_log$sticsa\_total)  
  
# next, subtract this average from all participants' sticsa scores,  
# and save as a variable  
df\_long\_ext\_fix\_duration\_log$sticsa\_total\_centred <-   
 df\_long\_ext\_fix\_duration\_log$sticsa\_total - df\_long\_ext\_fix\_duration\_log$sticsa\_total\_avg   
# from this we have mean sticsa scores after centring   
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) x 2 (Time: Early, Late) mixed ANOVA,   
# with mean-centred STICSA as covariate,  
# and obtain effect size (partial eta squared)  
ext\_fix\_duration\_ancova\_log <-   
 anova\_test(df\_long\_ext\_fix\_duration\_log,   
 fix\_duration\_log ~ iu\_group \* stimulus \* time + Error(id/(stimulus\*time)),  
 covariate = sticsa\_total\_centred, effect.size = "pes")   
  
# obtain the mixed ANCOVA results  
get\_anova\_table(ext\_fix\_duration\_ancova\_log)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 sticsa\_total\_centred 1 136 0.001 0.972 0.00000901  
## 2 iu\_group 1 136 8.365 0.004 \* 0.05800000  
## 3 stimulus 1 136 0.514 0.475 0.00400000  
## 4 time 1 136 4.358 0.039 \* 0.03100000  
## 5 sticsa\_total\_centred:stimulus 1 136 0.195 0.659 0.00100000  
## 6 iu\_group:stimulus 1 136 5.357 0.022 \* 0.03800000  
## 7 sticsa\_total\_centred:time 1 136 0.329 0.567 0.00200000  
## 8 iu\_group:time 1 136 0.501 0.480 0.00400000  
## 9 stimulus:time 1 136 0.174 0.677 0.00100000  
## 10 sticsa\_total\_centred:stimulus:time 1 136 0.221 0.639 0.00200000  
## 11 iu\_group:stimulus:time 1 136 0.379 0.539 0.00300000

# results:  
# STICSA (centred): F(1,136) = 0.01, p = .972, eta2(partial) < .001  
# IU: F(1,136) = 8.37, p = .004\*\*, eta2(partial) = .058  
# Stimulus: F(1,136) = 0.51, p = .475, eta2(partial) = .004  
# Time: F(1,136) = 4.36, p = .039\*, eta2(partial) = .031  
# STICSA \* Stimulus: F(1,136) = 0.20, p = .659, eta2(partial) = .001  
# IU \* Stimulus: F(1, 136) = 5.36, p = .022\*, eta2(partial) = .038  
# STICSA\* Time: F(1,136) = 0.33, p = .567, eta2(partial) = .002  
# IU \* Time: F(1,136) = 0.50, p = .480, eta2(partial) = . 004  
# Stimulus \* Time: F(1,136) = 0.17, p = .677, eta2(partial) = .001  
# STICSA \* Stimulus \* Time: F(1,136) = 0.22, p = .639, eta2(partial) = .002  
# IU \* Stimulus \* Time: F(1,136) = 0.34, p = .539, eta2(partial) = .003  
  
# there were significant main effects of IU, time,  
# and a significant IU-stimulus interaction on fixation duration in extinction,  
# and no further main effects or interactions.   
  
# write to csv  
write.csv((get\_anova\_table(ext\_fix\_duration\_ancova\_log)),   
 file = "tables/ancovas/ext\_fix\_duration\_ancova\_log.csv")  
  
# as there was a significant IU\*Stimulus interaction, conduct simple   
# main effects analysis:  
## obtain effect of IU at each level of stimulus   
simple\_effects\_ext\_fix\_duration\_iu\_ancova <- df\_long\_ext\_fix\_duration\_log %>%  
 group\_by(stimulus) %>%  
 anova\_test(dv = fix\_duration\_log, wid = id, between = iu\_group, within = time,  
 covariate = sticsa\_total\_centred, effect.size = "pes") %>%  
 get\_anova\_table() %>%  
 adjust\_pvalue(method = "bonferroni")  
  
# get the output  
simple\_effects\_ext\_fix\_duration\_iu\_ancova

## # A tibble: 10 × 9  
## stimulus Effect DFn DFd F p `p<.05` pes p.adj  
## \* <fct> <chr> <dbl> <dbl> <dbl> <dbl> <chr> <dbl> <dbl>  
## 1 -1 sticsa\_total\_centred 1 136 0.008 0.928 "" 6.02e-5 1   
## 2 -1 iu\_group 1 136 11.2 0.001 "\*" 7.6 e-2 0.01  
## 3 -1 time 1 136 4.37 0.038 "\*" 3.1 e-2 0.38  
## 4 -1 sticsa\_total\_centre… 1 136 0.627 0.43 "" 5 e-3 1   
## 5 -1 iu\_group:time 1 136 0.061 0.806 "" 4.45e-4 1   
## 6 1 sticsa\_total\_centred 1 136 0.027 0.87 "" 1.99e-4 1   
## 7 1 iu\_group 1 136 4.70 0.032 "\*" 3.3 e-2 0.32  
## 8 1 time 1 136 1.93 0.167 "" 1.4 e-2 1   
## 9 1 sticsa\_total\_centre… 1 136 0.036 0.849 "" 2.66e-4 1   
## 10 1 iu\_group:time 1 136 0.771 0.381 "" 6 e-3 1

# results:   
# The effect of IU group on CS+ was not significant [F(1,136) = 4.70, p = .320, pes = .033]  
# the effect of IU group on CS- was significant [F(1,136) = 11,19, p = .01, pes = .076]

## *ANCOVA Extinction Saccade Amplitude*

# transform wide format data into long format for mixed ANCOVA   
df\_long\_ext\_sacc\_amplitude <- melt(df, id = c("id", "iu\_group", "sticsa\_total"),   
 measure.vars = c("e\_ext\_csp\_sacc\_amplitude",   
 "e\_ext\_csm\_sacc\_amplitude",  
 "l\_ext\_csp\_sacc\_amplitude",   
 "l\_ext\_csm\_sacc\_amplitude"))  
  
# rename columns for easier interpretation  
colnames(df\_long\_ext\_sacc\_amplitude) = c("id", "iu\_group", "sticsa\_total", "condition", "sacc\_amplitude")  
  
# create column to code stimulus as CS+ (1) and CS- (-1)  
df\_long\_ext\_sacc\_amplitude$stimulus <-   
 factor(ifelse(df\_long\_ext\_sacc\_amplitude$condition == "e\_ext\_csp\_sacc\_amplitude" |   
 df\_long\_ext\_sacc\_amplitude$condition == "l\_ext\_csp\_sacc\_amplitude", 1, -1))  
  
# create column to code extinction as early (1) and late (-1)  
df\_long\_ext\_sacc\_amplitude$time <-   
 factor(ifelse(df\_long\_ext\_sacc\_amplitude$condition == "e\_ext\_csp\_sacc\_amplitude" |   
 df\_long\_ext\_sacc\_amplitude$condition == "e\_ext\_csm\_sacc\_amplitude", 1, -1))  
  
# mean centre continuous covariate (STICSA)  
# to apply mean centring, first obtain average sticsa scores for all participants,   
# and save as a variable  
df\_long\_ext\_sacc\_amplitude$sticsa\_total\_avg <- mean(df\_long\_ext\_sacc\_amplitude$sticsa\_total)  
  
# next, subtract this average from all participants' sticsa scores,  
# and save as a variable  
df\_long\_ext\_sacc\_amplitude$sticsa\_total\_centred <-   
 df\_long\_ext\_sacc\_amplitude$sticsa\_total - df\_long\_ext\_sacc\_amplitude$sticsa\_total\_avg   
# from this we have mean sticsa scores after centring   
  
# compute 2(IU: High & Low) x 2 (Stimulus: CS+, CS-) x 2 (Time: Early, Late) mixed ANOVA,   
# with mean-centred STICSA as covariate,  
# and obtain effect size (partial eta squared)  
ext\_sacc\_amplitude\_ancova <-   
 anova\_test(df\_long\_ext\_sacc\_amplitude,   
 sacc\_amplitude ~ iu\_group \* stimulus \* time + Error(id/(stimulus\*time)),  
 covariate = sticsa\_total\_centred, effect.size = "pes")

## Warning: NA detected in rows: 116,181,301.  
## Removing this rows before the analysis.

# obtain the mixed ANCOVA results  
get\_anova\_table(ext\_sacc\_amplitude\_ancova)

## ANOVA Table (type III tests)  
##   
## Effect DFn DFd F p p<.05 pes  
## 1 sticsa\_total\_centred 1 133 1.134 0.289 0.008000  
## 2 iu\_group 1 133 1.025 0.313 0.008000  
## 3 stimulus 1 133 0.754 0.387 0.006000  
## 4 time 1 133 0.255 0.615 0.002000  
## 5 sticsa\_total\_centred:stimulus 1 133 0.370 0.544 0.003000  
## 6 iu\_group:stimulus 1 133 2.035 0.156 0.015000  
## 7 sticsa\_total\_centred:time 1 133 1.359 0.246 0.010000  
## 8 iu\_group:time 1 133 0.803 0.372 0.006000  
## 9 stimulus:time 1 133 0.071 0.790 0.000533  
## 10 sticsa\_total\_centred:stimulus:time 1 133 0.421 0.517 0.003000  
## 11 iu\_group:stimulus:time 1 133 0.997 0.320 0.007000

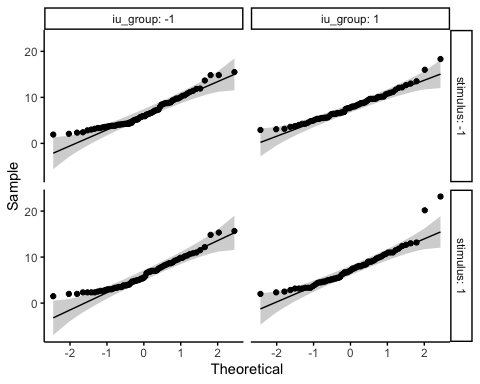
# results:  
# STICSA (centred): F(1,133) = 1.13, p = .289, eta2(partial) = .008  
# IU: F(1,133) = 1.03, p = .313, eta2(partial) = .008  
# Stimulus: F(1,133) = 0.75, p = .387, eta2(partial) = .006  
# Time: F(1,133) = 0.26, p = .615, eta2(partial) = .002  
# STICSA \* Stimulus: F(1,133) = 0.37, p = .544, eta2(partial) = .003  
# IU \* Stimulus: F(1, 133) = 2.04, p = .156, eta2(partial) = .015  
# STICSA\* Time: F(1,133) = 1.36, p = .246, eta2(partial) = .010  
# IU \* Time: F(1,133) = 0.80, p = .372, eta2(partial) = . 006  
# Stimulus \* Time: F(1,133) = 0.07, p = .790, eta2(partial) = .001  
# STICSA \* Stimulus \* Time: F(1,133) = 0.42, p = .517, eta2(partial) = .003  
# IU \* Stimulus \* Time: F(1,133) = 0.10, p = .320, eta2(partial) = .007  
  
# therefore, even after accounting for trait anxiety, there continue  
# to be no significant effects or interactions on saccade amplitude  
# in extinction  
  
# write to csv  
write.csv((get\_anova\_table(ext\_sacc\_amplitude\_ancova)),   
 file = "tables/ancovas/ext\_sacc\_amplitude\_ancova.csv")

# *Assumption Checks*

############ assumptions of mixed ANOVA:  
# categorical IVs, interval/ratio DVs  
# outcome variable(s) should be approximately normally distributed  
# no significant outliers in the groups  
# homogeneity of variances  
# sphericity (not applicable in this case, as no within-subjects factors with > 3 levels)  
# homogeneity of variance-covariance matrices   
  
############ additional assumptions of ANCOVA:  
# independence of covariate and IVs  
# homogeneity of regression slopes  
# linearity between covariate and outcome variable(s) at each level of grouping variables

## *Normality of Outcome Variables*

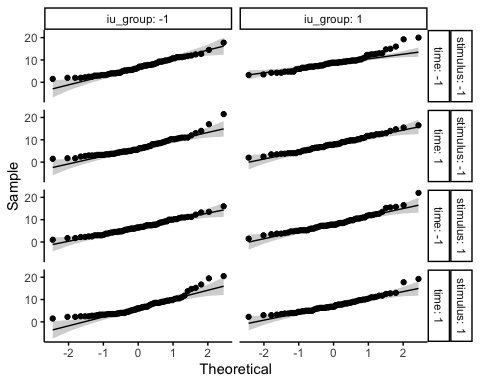
############### note: variables coded as follows:  
#### IU  
# high IU: 1  
# low IU: -1  
  
#### stimulus  
# CS+: 1  
# CS-: -1  
  
#### time  
# early: 1  
# late: -1  
  
######### acquisition fix count  
## check QQ plot  
qqplot\_acq\_fix\_count <- ggqqplot(df\_long\_acq\_fix\_count, "fix\_count", ggtheme = theme\_classic()) +  
 facet\_grid(stimulus ~ iu\_group, labeller = "label\_both")  
qqplot\_acq\_fix\_count



## check shapiro  
shapiro\_acq\_fix\_count <- df\_long\_acq\_fix\_count %>%  
 group\_by(iu\_group, stimulus) %>%  
 shapiro\_test(fix\_count)  
shapiro\_acq\_fix\_count

## # A tibble: 4 × 5  
## iu\_group stimulus variable statistic p  
## <fct> <fct> <chr> <dbl> <dbl>  
## 1 -1 -1 fix\_count 0.929 0.000611   
## 2 -1 1 fix\_count 0.934 0.00108   
## 3 1 -1 fix\_count 0.962 0.0364   
## 4 1 1 fix\_count 0.895 0.0000312

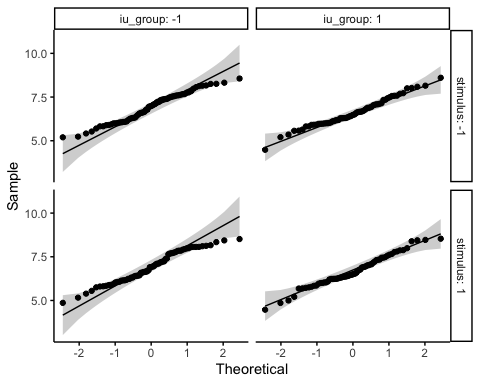
# p-values < .05: data violate assumption of normality  
  
  
######### extinction fix count  
## check QQ plot  
qqplot\_ext\_fix\_count <- ggqqplot(df\_long\_ext\_fix\_count, "fix\_count", ggtheme = theme\_classic()) +  
 facet\_grid(stimulus + time ~ iu\_group, labeller = "label\_both")  
qqplot\_ext\_fix\_count



## check shapiro  
shapiro\_ext\_fix\_count <-df\_long\_ext\_fix\_count %>%  
 group\_by(iu\_group, stimulus, time) %>%  
 shapiro\_test(fix\_count)  
shapiro\_ext\_fix\_count

## # A tibble: 8 × 6  
## iu\_group stimulus time variable statistic p  
## <fct> <fct> <fct> <chr> <dbl> <dbl>  
## 1 -1 -1 -1 fix\_count 0.961 0.0263   
## 2 -1 -1 1 fix\_count 0.904 0.0000488   
## 3 -1 1 -1 fix\_count 0.977 0.228   
## 4 -1 1 1 fix\_count 0.881 0.00000681  
## 5 1 -1 -1 fix\_count 0.929 0.000810   
## 6 1 -1 1 fix\_count 0.981 0.391   
## 7 1 1 -1 fix\_count 0.945 0.00457   
## 8 1 1 1 fix\_count 0.931 0.000995

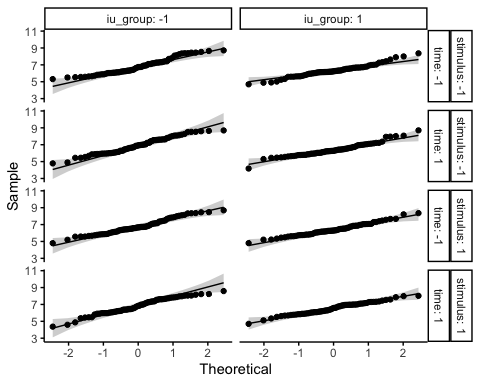
# p-values < .05: data violate assumption of normality for all except:  
# high IU late extinction CS- and low IU early extinction CS+ (ps > .05)  
  
######### acquisition fix duration log  
## check QQ plot  
qqplot\_acq\_fix\_duration\_log <- ggqqplot(df\_long\_acq\_fix\_duration\_log, "fix\_duration\_log", ggtheme = theme\_classic()) +  
 facet\_grid(stimulus ~ iu\_group, labeller = "label\_both")  
qqplot\_acq\_fix\_duration\_log



## check shapiro  
shapiro\_acq\_fix\_duration\_log <- df\_long\_acq\_fix\_duration\_log %>%  
 group\_by(iu\_group, stimulus) %>%  
 shapiro\_test(fix\_duration\_log)  
shapiro\_acq\_fix\_duration\_log

## # A tibble: 4 × 5  
## iu\_group stimulus variable statistic p  
## <fct> <fct> <chr> <dbl> <dbl>  
## 1 -1 -1 fix\_duration\_log 0.970 0.0814  
## 2 -1 1 fix\_duration\_log 0.964 0.0398  
## 3 1 -1 fix\_duration\_log 0.981 0.385   
## 4 1 1 fix\_duration\_log 0.981 0.408

# p-values > .05: data meet assumption of normality for all except:  
# low IU CS+ (p = .039)  
  
######### extinction fix duration log  
## check QQ plot  
qqplot\_ext\_fix\_duration\_log <- ggqqplot(df\_long\_ext\_fix\_duration\_log, "fix\_duration\_log", ggtheme = theme\_classic()) +  
 facet\_grid(stimulus + time ~ iu\_group, labeller = "label\_both")  
qqplot\_ext\_fix\_duration\_log



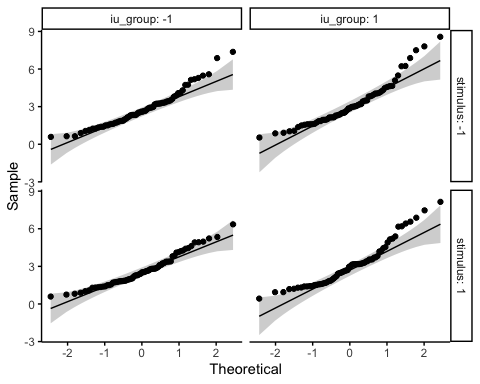
## check shapiro  
shapiro\_ext\_fix\_duration\_log <- df\_long\_ext\_fix\_duration\_log %>%  
 group\_by(iu\_group, stimulus, time) %>%  
 shapiro\_test(fix\_duration\_log)  
shapiro\_ext\_fix\_duration\_log

## # A tibble: 8 × 6  
## iu\_group stimulus time variable statistic p  
## <fct> <fct> <fct> <chr> <dbl> <dbl>  
## 1 -1 -1 -1 fix\_duration\_log 0.945 0.00364  
## 2 -1 -1 1 fix\_duration\_log 0.974 0.143   
## 3 -1 1 -1 fix\_duration\_log 0.972 0.112   
## 4 -1 1 1 fix\_duration\_log 0.970 0.0913   
## 5 1 -1 -1 fix\_duration\_log 0.973 0.146   
## 6 1 -1 1 fix\_duration\_log 0.959 0.0242   
## 7 1 1 -1 fix\_duration\_log 0.984 0.523   
## 8 1 1 1 fix\_duration\_log 0.983 0.460

# p-values > .05: data meet assumption of normality for all except:  
# low IU CS- early extinction and high IU CS- late extinction  
  
######### acquisition sacc amplitude  
## check QQ plot  
qqplot\_acq\_sacc\_amplitude <- ggqqplot(df\_long\_acq\_sacc\_amplitude, "sacc\_amplitude", ggtheme = theme\_classic()) +  
 facet\_grid(stimulus ~ iu\_group, labeller = "label\_both")  
qqplot\_acq\_sacc\_amplitude

## Warning: Removed 2 rows containing non-finite values (stat\_qq).

## Warning: Removed 2 rows containing non-finite values (stat\_qq\_line).  
  
## Warning: Removed 2 rows containing non-finite values (stat\_qq\_line).



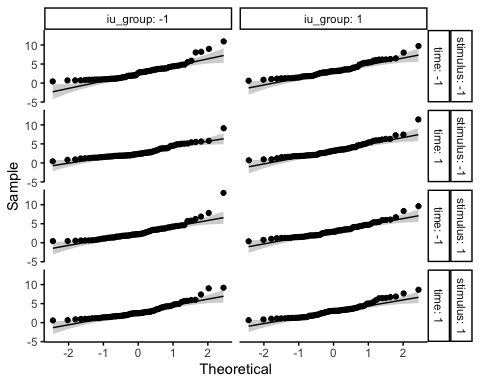
## check shapiro  
shapiro\_acq\_sacc\_amplitude <- df\_long\_acq\_sacc\_amplitude %>%  
 group\_by(iu\_group, stimulus) %>%  
 shapiro\_test(sacc\_amplitude)  
shapiro\_acq\_sacc\_amplitude

## # A tibble: 4 × 5  
## iu\_group stimulus variable statistic p  
## <fct> <fct> <chr> <dbl> <dbl>  
## 1 -1 -1 sacc\_amplitude 0.940 0.00227   
## 2 -1 1 sacc\_amplitude 0.954 0.0111   
## 3 1 -1 sacc\_amplitude 0.913 0.000176  
## 4 1 1 sacc\_amplitude 0.918 0.000275

# p-values < .05: data violate assumption of normality  
  
######### extinction sacc amplitude  
## check QQ plot  
qqplot\_ext\_sacc\_amplitude <- ggqqplot(df\_long\_ext\_sacc\_amplitude, "sacc\_amplitude", ggtheme = theme\_classic()) +  
 facet\_grid(stimulus + time ~ iu\_group, labeller = "label\_both")  
qqplot\_ext\_sacc\_amplitude

## Warning: Removed 3 rows containing non-finite values (stat\_qq).

## Warning: Removed 3 rows containing non-finite values (stat\_qq\_line).  
  
## Warning: Removed 3 rows containing non-finite values (stat\_qq\_line).



## check shapiro  
shapiro\_ext\_sacc\_amplitude <- df\_long\_ext\_sacc\_amplitude %>%  
 group\_by(iu\_group, stimulus, time) %>%  
 shapiro\_test(sacc\_amplitude)  
shapiro\_ext\_sacc\_amplitude

## # A tibble: 8 × 6  
## iu\_group stimulus time variable statistic p  
## <fct> <fct> <fct> <chr> <dbl> <dbl>  
## 1 -1 -1 -1 sacc\_amplitude 0.849 0.000000535   
## 2 -1 -1 1 sacc\_amplitude 0.889 0.0000125   
## 3 -1 1 -1 sacc\_amplitude 0.821 0.0000000925  
## 4 -1 1 1 sacc\_amplitude 0.880 0.00000688   
## 5 1 -1 -1 sacc\_amplitude 0.930 0.000946   
## 6 1 -1 1 sacc\_amplitude 0.902 0.0000659   
## 7 1 1 -1 sacc\_amplitude 0.925 0.000514   
## 8 1 1 1 sacc\_amplitude 0.926 0.000578

# p-values < .05: data violate assumption of normality

## *Outliers*

# identify outliers using identify\_outliers function from rstatix package,  
# where third quartile + 3xIQR or below first quartile - 3xIQR   
# are considered as extreme points (or extreme outliers).  
  
## acquisition fix count  
outliers\_acq\_fix\_count <- df\_long\_acq\_fix\_count %>%  
 group\_by(iu\_group, stimulus) %>%  
 identify\_outliers(fix\_count)  
outliers\_acq\_fix\_count

## # A tibble: 4 × 10  
## iu\_group stimulus id sticsa\_total condition fix\_count sticsa\_total\_avg  
## <fct> <fct> <fct> <dbl> <fct> <dbl> <dbl>  
## 1 1 -1 086\_1 68 acq\_csm\_fix\_c… 18.3 40.5  
## 2 1 -1 099\_1 52 acq\_csm\_fix\_c… 16 40.5  
## 3 1 1 086\_1 68 acq\_csp\_fix\_c… 23.2 40.5  
## 4 1 1 099\_1 52 acq\_csp\_fix\_c… 20.2 40.5  
## # … with 3 more variables: sticsa\_total\_centred <dbl>, is.outlier <lgl>,  
## # is.extreme <lgl>

# no extreme outliers  
  
## extinction fix count  
outliers\_ext\_fix\_count <-   
df\_long\_ext\_fix\_count %>%  
 group\_by(iu\_group, stimulus, time) %>%  
 identify\_outliers(fix\_count)  
outliers\_ext\_fix\_count

## # A tibble: 14 × 11  
## iu\_group stimulus time id sticsa\_total condition fix\_count  
## <fct> <fct> <fct> <fct> <dbl> <fct> <dbl>  
## 1 -1 -1 -1 122\_1 37 l\_ext\_csm\_fix\_count 17.8  
## 2 -1 -1 1 047\_1 41 e\_ext\_csm\_fix\_count 17   
## 3 -1 -1 1 122\_1 37 e\_ext\_csm\_fix\_count 21.5  
## 4 -1 1 -1 122\_1 37 l\_ext\_csp\_fix\_count 16   
## 5 -1 1 1 122\_1 37 e\_ext\_csp\_fix\_count 20.5  
## 6 -1 1 1 143\_1 44 e\_ext\_csp\_fix\_count 19.5  
## 7 1 -1 -1 033\_1 54 l\_ext\_csm\_fix\_count 20   
## 8 1 -1 -1 065\_1 33 l\_ext\_csm\_fix\_count 14.8  
## 9 1 -1 -1 086\_1 68 l\_ext\_csm\_fix\_count 19.2  
## 10 1 -1 -1 099\_1 52 l\_ext\_csm\_fix\_count 16   
## 11 1 -1 -1 113\_1 31 l\_ext\_csm\_fix\_count 15   
## 12 1 1 -1 086\_1 68 l\_ext\_csp\_fix\_count 22   
## 13 1 1 1 086\_1 68 e\_ext\_csp\_fix\_count 19.2  
## 14 1 1 1 113\_1 31 e\_ext\_csp\_fix\_count 17.8  
## # … with 4 more variables: sticsa\_total\_avg <dbl>, sticsa\_total\_centred <dbl>,  
## # is.outlier <lgl>, is.extreme <lgl>

# two extreme outliers: ppt 33 and 86  
  
# acquisition fix duration log  
outliers\_acq\_fix\_duration\_log <- df\_long\_acq\_fix\_duration\_log %>%  
 group\_by(iu\_group, stimulus) %>%  
 identify\_outliers(fix\_duration\_log)  
# no extreme outliers  
  
outliers\_acq\_fix\_duration\_log

## [1] iu\_group stimulus id   
## [4] sticsa\_total condition fix\_duration\_log   
## [7] sticsa\_total\_avg sticsa\_total\_centred is.outlier   
## [10] is.extreme   
## <0 rows> (or 0-length row.names)

## extinction fix duration log  
outliers\_ext\_fix\_duration\_log <- df\_long\_ext\_fix\_duration\_log %>%  
 group\_by(iu\_group, stimulus, time) %>%  
 identify\_outliers(fix\_duration\_log)  
outliers\_ext\_fix\_duration\_log

## # A tibble: 6 × 11  
## iu\_group stimulus time id sticsa\_total condition fix\_duration\_log  
## <fct> <fct> <fct> <fct> <dbl> <fct> <dbl>  
## 1 1 -1 -1 009\_1 41 l\_ext\_csm\_fix\_dur… 8.00  
## 2 1 -1 -1 010\_1 43 l\_ext\_csm\_fix\_dur… 4.69  
## 3 1 -1 -1 015\_1 55 l\_ext\_csm\_fix\_dur… 8.37  
## 4 1 -1 -1 044\_1 36 l\_ext\_csm\_fix\_dur… 7.91  
## 5 1 -1 1 044\_1 36 e\_ext\_csm\_fix\_dur… 8.70  
## 6 1 -1 1 113\_1 31 e\_ext\_csm\_fix\_dur… 4.18  
## # … with 4 more variables: sticsa\_total\_avg <dbl>, sticsa\_total\_centred <dbl>,  
## # is.outlier <lgl>, is.extreme <lgl>

# no extreme outliers  
  
## acquisition sacc amplitude  
outliers\_acq\_sacc\_amplitude <- df\_long\_acq\_sacc\_amplitude %>%  
 group\_by(iu\_group, stimulus) %>%  
 identify\_outliers(sacc\_amplitude)  
outliers\_acq\_sacc\_amplitude

## # A tibble: 9 × 10  
## iu\_group stimulus id sticsa\_total condition sacc\_amplitude sticsa\_total\_avg  
## <fct> <fct> <fct> <dbl> <fct> <dbl> <dbl>  
## 1 -1 -1 016\_1 26 acq\_csm\_… 7.37 40.5  
## 2 -1 -1 026\_1 55 acq\_csm\_… 6.87 40.5  
## 3 -1 1 016\_1 26 acq\_csp\_… 6.35 40.5  
## 4 1 -1 017\_1 33 acq\_csm\_… 7.81 40.5  
## 5 1 -1 021\_1 54 acq\_csm\_… 7.50 40.5  
## 6 1 -1 022\_1 50 acq\_csm\_… 8.57 40.5  
## 7 1 1 009\_1 41 acq\_csp\_… 7.47 40.5  
## 8 1 1 043\_1 39 acq\_csp\_… 6.88 40.5  
## 9 1 1 044\_1 36 acq\_csp\_… 8.15 40.5  
## # … with 3 more variables: sticsa\_total\_centred <dbl>, is.outlier <lgl>,  
## # is.extreme <lgl>

# no extreme outliers  
  
## extinction sacc amplitude  
outliers\_ext\_sacc\_amplitude <- df\_long\_ext\_sacc\_amplitude %>%  
 group\_by(iu\_group, stimulus, time) %>%  
 identify\_outliers(sacc\_amplitude)  
outliers\_ext\_sacc\_amplitude

## # A tibble: 17 × 11  
## iu\_group stimulus time id sticsa\_total condition sacc\_amplitude  
## <fct> <fct> <fct> <fct> <dbl> <fct> <dbl>  
## 1 -1 -1 -1 016\_1 26 l\_ext\_csm\_sacc\_amp… 10.9   
## 2 -1 -1 -1 075\_1 35 l\_ext\_csm\_sacc\_amp… 8.98  
## 3 -1 -1 -1 078\_1 42 l\_ext\_csm\_sacc\_amp… 8.03  
## 4 -1 -1 -1 111\_1 41 l\_ext\_csm\_sacc\_amp… 8.21  
## 5 -1 -1 1 016\_1 26 e\_ext\_csm\_sacc\_amp… 9.11  
## 6 -1 1 -1 016\_1 26 l\_ext\_csp\_sacc\_amp… 13.1   
## 7 -1 1 -1 075\_1 35 l\_ext\_csp\_sacc\_amp… 7.84  
## 8 -1 1 1 016\_1 26 e\_ext\_csp\_sacc\_amp… 9.02  
## 9 -1 1 1 051\_1 28 e\_ext\_csp\_sacc\_amp… 7.40  
## 10 -1 1 1 119\_1 43 e\_ext\_csp\_sacc\_amp… 9.18  
## 11 1 -1 -1 009\_1 41 l\_ext\_csm\_sacc\_amp… 8.00  
## 12 1 -1 -1 105\_1 33 l\_ext\_csm\_sacc\_amp… 9.74  
## 13 1 -1 1 105\_1 33 e\_ext\_csm\_sacc\_amp… 11.4   
## 14 1 1 -1 009\_1 41 l\_ext\_csp\_sacc\_amp… 9.62  
## 15 1 1 -1 022\_1 50 l\_ext\_csp\_sacc\_amp… 8.34  
## 16 1 1 1 009\_1 41 e\_ext\_csp\_sacc\_amp… 7.67  
## 17 1 1 1 129\_1 46 e\_ext\_csp\_sacc\_amp… 8.65  
## # … with 4 more variables: sticsa\_total\_avg <dbl>, sticsa\_total\_centred <dbl>,  
## # is.outlier <lgl>, is.extreme <lgl>

# two extreme outliers: ppt 16 and 105

## *Homogeneity of Variance*

# this will be done using levene's test  
  
## acquisition fix count  
levene\_acq\_fix\_count <- df\_long\_acq\_fix\_count %>%  
 group\_by(stimulus) %>%  
 levene\_test(fix\_count ~ iu\_group)  
levene\_acq\_fix\_count

## # A tibble: 2 × 5  
## stimulus df1 df2 statistic p  
## <fct> <int> <int> <dbl> <dbl>  
## 1 -1 1 137 0.477 0.491  
## 2 1 1 137 0.0415 0.839

# p-values > .05, data meet assumption of homogeneity of variance  
  
## extinction fix count  
levene\_ext\_fix\_count <-df\_long\_ext\_fix\_count %>%  
 group\_by(stimulus, time) %>%  
 levene\_test(fix\_count ~ iu\_group)  
levene\_ext\_fix\_count

## # A tibble: 4 × 6  
## stimulus time df1 df2 statistic p  
## <fct> <fct> <int> <int> <dbl> <dbl>  
## 1 -1 -1 1 137 1.45 0.231  
## 2 -1 1 1 137 0.181 0.671  
## 3 1 -1 1 137 0.264 0.608  
## 4 1 1 1 137 1.86 0.174

# p-values > .05, data meet assumption of homogeneity of variance  
  
# acquisition fix duration log  
levene\_acq\_fix\_duration\_log <- df\_long\_acq\_fix\_duration\_log %>%  
 group\_by(stimulus) %>%  
 levene\_test(fix\_duration\_log ~ iu\_group)  
levene\_acq\_fix\_duration\_log

## # A tibble: 2 × 5  
## stimulus df1 df2 statistic p  
## <fct> <int> <int> <dbl> <dbl>  
## 1 -1 1 137 2.04 0.155  
## 2 1 1 137 0.753 0.387

# p-values > .05, data meet assumption of homogeneity of variance  
  
## extinction fix count  
levene\_ext\_fix\_duration\_log <- df\_long\_ext\_fix\_duration\_log %>%  
 group\_by(stimulus, time) %>%  
 levene\_test(fix\_duration\_log ~ iu\_group)  
levene\_ext\_fix\_duration\_log

## # A tibble: 4 × 6  
## stimulus time df1 df2 statistic p  
## <fct> <fct> <int> <int> <dbl> <dbl>  
## 1 -1 -1 1 137 8.18 0.00490  
## 2 -1 1 1 137 7.74 0.00616  
## 3 1 -1 1 137 2.78 0.0977   
## 4 1 1 1 137 7.14 0.00843

# p-value for early extinction and CS+ > .05, data meet assumption of homogeneity of variance  
# p-values for early extinction and CS-, and late extinction and both stimulli < .05,   
# data violate assumption of homogeneity of variance  
  
## acquisition sacc amplitude  
levene\_acq\_sacc\_amplitude <- df\_long\_acq\_sacc\_amplitude %>%  
 group\_by(stimulus) %>%  
 levene\_test(sacc\_amplitude ~ iu\_group)  
levene\_acq\_sacc\_amplitude

## # A tibble: 2 × 5  
## stimulus df1 df2 statistic p  
## <fct> <int> <int> <dbl> <dbl>  
## 1 -1 1 135 1.03 0.311   
## 2 1 1 137 3.42 0.0665

# p-values > .05, data meet assumption of homogeneity of variance  
  
## extinction sacc amplitude  
levene\_ext\_sacc\_amplitude <- df\_long\_ext\_sacc\_amplitude %>%  
 group\_by(stimulus, time) %>%  
 levene\_test(sacc\_amplitude ~ iu\_group)  
levene\_ext\_sacc\_amplitude

## # A tibble: 4 × 6  
## stimulus time df1 df2 statistic p  
## <fct> <fct> <int> <int> <dbl> <dbl>  
## 1 -1 -1 1 137 0.364 0.547  
## 2 -1 1 1 136 1.72 0.191  
## 3 1 -1 1 136 0.0230 0.880  
## 4 1 1 1 136 0.0324 0.857

# p-values > .05, data meet assumption of homogeneity of variance  
  
# however, in large samples, levene's test can be sig even when group variances  
# are not very different.

## *Homogeneity of Variance-Covariance Matrices*

# this tests whether covariance matrices are equal across cells formed by  
# between-subjects factor (IU)  
  
# use Box's M (however, this is highly sensitive, so unless p < .001 and sample  
# sizes are unequal, can ignore it)  
  
box\_m\_acq\_fix\_count <-   
box\_m(df\_long\_acq\_fix\_count[, "fix\_count", drop = FALSE], df\_long\_acq\_fix\_count$iu\_group)  
box\_m\_acq\_fix\_count

## # A tibble: 1 × 4  
## statistic p.value parameter method   
## <dbl> <dbl> <dbl> <chr>   
## 1 0.224 0.636 1 Box's M-test for Homogeneity of Covariance Matric…

# p-value > .05, data meet assumption of homogeneity of variance-covariance matrices  
  
box\_m\_ext\_fix\_count <-   
box\_m(df\_long\_ext\_fix\_count[, "fix\_count", drop = FALSE], df\_long\_ext\_fix\_count$iu\_group)  
box\_m\_ext\_fix\_count

## # A tibble: 1 × 4  
## statistic p.value parameter method   
## <dbl> <dbl> <dbl> <chr>   
## 1 0.753 0.385 1 Box's M-test for Homogeneity of Covariance Matric…

# p-value > .05, data meet assumption of homogeneity of variance-covariance matrices  
  
bom\_m\_acq\_fix\_duration\_log <-  
box\_m(df\_long\_acq\_fix\_duration\_log[, "fix\_duration\_log", drop = FALSE], df\_long\_acq\_fix\_duration\_log$iu\_group)  
bom\_m\_acq\_fix\_duration\_log

## # A tibble: 1 × 4  
## statistic p.value parameter method   
## <dbl> <dbl> <dbl> <chr>   
## 1 0.358 0.550 1 Box's M-test for Homogeneity of Covariance Matric…

# p-value > .05, data meet assumption of homogeneity of variance-covariance matrices  
  
box\_m\_ext\_fix\_duration\_log <-  
box\_m(df\_long\_ext\_fix\_duration\_log[, "fix\_duration\_log", drop = FALSE], df\_long\_ext\_fix\_duration\_log$iu\_group)  
box\_m\_ext\_fix\_duration\_log

## # A tibble: 1 × 4  
## statistic p.value parameter method   
## <dbl> <dbl> <dbl> <chr>   
## 1 16.7 0.0000435 1 Box's M-test for Homogeneity of Covariance Matr…

# p-value < .05, data violate assumption of homogeneity of variance-covariance matrices

## *Independence of Covariate and IVs*

### *Fixation Count*

#### *Acquisition*

# sticsa and iu group  
t\_test\_independence\_sticsa\_iu\_group\_acq\_fix\_count <-   
 t.test(  
 df\_long\_acq\_fix\_count[df\_long\_acq\_fix\_count$iu\_group == "1", "sticsa\_total\_centred"],  
 df\_long\_acq\_fix\_count[df\_long\_acq\_fix\_count$iu\_group == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_iu\_group\_acq\_fix\_count

##   
## Two Sample t-test  
##   
## data: df\_long\_acq\_fix\_count[df\_long\_acq\_fix\_count$iu\_group == "1", "sticsa\_total\_centred"] and df\_long\_acq\_fix\_count[df\_long\_acq\_fix\_count$iu\_group == "-1", "sticsa\_total\_centred"]  
## t = 9.3255, df = 276, p-value < 0.00000000000000022  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## 7.343247 11.273157  
## sample estimates:  
## mean of x mean of y   
## 4.754549 -4.553653

# p < .05 : sticsa is not independent of iu group   
  
# sticsa and stimulus  
t\_test\_independence\_sticsa\_stimulus\_acq\_fix\_count <-   
 t.test(  
 df\_long\_acq\_fix\_count[df\_long\_acq\_fix\_count$stimulus == "1", "sticsa\_total\_centred"],  
 df\_long\_acq\_fix\_count[df\_long\_acq\_fix\_count$stimulus == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_stimulus\_acq\_fix\_count

##   
## Two Sample t-test  
##   
## data: df\_long\_acq\_fix\_count[df\_long\_acq\_fix\_count$stimulus == "1", "sticsa\_total\_centred"] and df\_long\_acq\_fix\_count[df\_long\_acq\_fix\_count$stimulus == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 276, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -2.252832 2.252832  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862893 -0.000000000000002862893

# p > .05 - sticsa is independent of stimulus

#### *Extinction*

# sticsa and iu group  
t\_test\_independence\_sticsa\_iu\_group\_ext\_fix\_count <-   
 t.test(  
 df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_iu\_group\_ext\_fix\_count

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "1", "sticsa\_total\_centred"] and df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$iu\_group == "-1", "sticsa\_total\_centred"]  
## t = 13.212, df = 554, p-value < 0.00000000000000022  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## 7.924338 10.692067  
## sample estimates:  
## mean of x mean of y   
## 4.754549 -4.553653

# p < .05 : sticsa is not independent of iu group   
  
# sticsa and stimulus  
t\_test\_independence\_sticsa\_stimulus\_ext\_fix\_count <-   
 t.test(  
 df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$stimulus == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$stimulus == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_stimulus\_ext\_fix\_count

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$stimulus == "1", "sticsa\_total\_centred"] and df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$stimulus == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 554, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -1.586608 1.586608  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862855 -0.000000000000002862855

# p > .05 - sticsa is independent of stimulus  
  
# sticsa and time  
t\_test\_independence\_sticsa\_time\_ext\_fix\_count <-   
 t.test(  
 df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$time == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$time == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_time\_ext\_fix\_count

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$time == "1", "sticsa\_total\_centred"] and df\_long\_ext\_fix\_count[df\_long\_ext\_fix\_count$time == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 554, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -1.586608 1.586608  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862855 -0.000000000000002862855

# p > .05 - sticsa is independent of time

### *Fixation Duration (Log Transformed)*

#### *Acquisition*

# sticsa and iu group  
t\_test\_independence\_sticsa\_iu\_group\_acq\_fix\_duration\_log <-   
 t.test(  
 df\_long\_acq\_fix\_duration\_log[df\_long\_acq\_fix\_duration\_log$iu\_group == "1", "sticsa\_total\_centred"],  
 df\_long\_acq\_fix\_duration\_log[df\_long\_acq\_fix\_duration\_log$iu\_group == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_iu\_group\_acq\_fix\_duration\_log

##   
## Two Sample t-test  
##   
## data: df\_long\_acq\_fix\_duration\_log[df\_long\_acq\_fix\_duration\_log$iu\_group == "1", "sticsa\_total\_centred"] and df\_long\_acq\_fix\_duration\_log[df\_long\_acq\_fix\_duration\_log$iu\_group == "-1", "sticsa\_total\_centred"]  
## t = 9.3255, df = 276, p-value < 0.00000000000000022  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## 7.343247 11.273157  
## sample estimates:  
## mean of x mean of y   
## 4.754549 -4.553653

# p < .05 : sticsa is not independent of iu group   
  
# sticsa and stimulus  
t\_test\_independence\_sticsa\_stimulus\_acq\_fix\_duration\_log <-   
 t.test(  
 df\_long\_acq\_fix\_duration\_log[df\_long\_acq\_fix\_duration\_log$stimulus == "1", "sticsa\_total\_centred"],  
 df\_long\_acq\_fix\_duration\_log[df\_long\_acq\_fix\_duration\_log$stimulus == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_stimulus\_acq\_fix\_duration\_log

##   
## Two Sample t-test  
##   
## data: df\_long\_acq\_fix\_duration\_log[df\_long\_acq\_fix\_duration\_log$stimulus == "1", "sticsa\_total\_centred"] and df\_long\_acq\_fix\_duration\_log[df\_long\_acq\_fix\_duration\_log$stimulus == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 276, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -2.252832 2.252832  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862893 -0.000000000000002862893

# p > .05 - sticsa is independent of stimulus

#### *Extinction*

# sticsa and iu group  
t\_test\_independence\_sticsa\_iu\_group\_ext\_fix\_duration\_log <-   
 t.test(  
 df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_iu\_group\_ext\_fix\_duration\_log

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "1", "sticsa\_total\_centred"] and df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$iu\_group == "-1", "sticsa\_total\_centred"]  
## t = 13.212, df = 554, p-value < 0.00000000000000022  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## 7.924338 10.692067  
## sample estimates:  
## mean of x mean of y   
## 4.754549 -4.553653

# p < .05 : sticsa is not independent of iu group   
  
# sticsa and stimulus  
t\_test\_independence\_sticsa\_stimulus\_ext\_fix\_duration\_log <-   
 t.test(  
 df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$stimulus == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$stimulus == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_stimulus\_ext\_fix\_duration\_log

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$stimulus == "1", "sticsa\_total\_centred"] and df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$stimulus == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 554, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -1.586608 1.586608  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862855 -0.000000000000002862855

# p > .05 - sticsa is independent of stimulus  
  
# sticsa and time  
t\_test\_independence\_sticsa\_time\_ext\_fix\_duration <-   
 t.test(  
 df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$time == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$time == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_time\_ext\_fix\_duration

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$time == "1", "sticsa\_total\_centred"] and df\_long\_ext\_fix\_duration\_log[df\_long\_ext\_fix\_duration\_log$time == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 554, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -1.586608 1.586608  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862855 -0.000000000000002862855

# p > .05 - sticsa is independent of time

### *Saccade Amplitude*

#### *Acquisition*

# sticsa and iu group  
t\_test\_independence\_sticsa\_iu\_group\_acq\_sacc\_amplitude <-   
 t.test(  
 df\_long\_acq\_sacc\_amplitude[df\_long\_acq\_sacc\_amplitude$iu\_group == "1", "sticsa\_total\_centred"],  
 df\_long\_acq\_sacc\_amplitude[df\_long\_acq\_sacc\_amplitude$iu\_group == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_iu\_group\_acq\_sacc\_amplitude

##   
## Two Sample t-test  
##   
## data: df\_long\_acq\_sacc\_amplitude[df\_long\_acq\_sacc\_amplitude$iu\_group == "1", "sticsa\_total\_centred"] and df\_long\_acq\_sacc\_amplitude[df\_long\_acq\_sacc\_amplitude$iu\_group == "-1", "sticsa\_total\_centred"]  
## t = 9.3255, df = 276, p-value < 0.00000000000000022  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## 7.343247 11.273157  
## sample estimates:  
## mean of x mean of y   
## 4.754549 -4.553653

# p < .05 : sticsa is not independent of iu group   
  
# sticsa and stimulus  
t\_test\_independence\_sticsa\_stimulus\_acq\_sacc\_amplitude <-   
 t.test(  
 df\_long\_acq\_sacc\_amplitude[df\_long\_acq\_sacc\_amplitude$stimulus == "1", "sticsa\_total\_centred"],  
 df\_long\_acq\_sacc\_amplitude[df\_long\_acq\_sacc\_amplitude$stimulus == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_stimulus\_acq\_sacc\_amplitude

##   
## Two Sample t-test  
##   
## data: df\_long\_acq\_sacc\_amplitude[df\_long\_acq\_sacc\_amplitude$stimulus == "1", "sticsa\_total\_centred"] and df\_long\_acq\_sacc\_amplitude[df\_long\_acq\_sacc\_amplitude$stimulus == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 276, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -2.252832 2.252832  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862893 -0.000000000000002862893

# p > .05 - sticsa is independent of stimulus

#### *Extinction*

# sticsa and iu group  
t\_test\_independence\_sticsa\_iu\_group\_ext\_sacc\_amplitude <-   
 t.test(  
 df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_iu\_group\_ext\_sacc\_amplitude

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "1", "sticsa\_total\_centred"] and df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$iu\_group == "-1", "sticsa\_total\_centred"]  
## t = 13.212, df = 554, p-value < 0.00000000000000022  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## 7.924338 10.692067  
## sample estimates:  
## mean of x mean of y   
## 4.754549 -4.553653

# p < .05 : sticsa is not independent of iu group   
  
# sticsa and stimulus  
t\_test\_independence\_sticsa\_stimulus\_ext\_sacc\_amplitude <-   
 t.test(  
 df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$stimulus == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$stimulus == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_stimulus\_ext\_sacc\_amplitude

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$stimulus == "1", "sticsa\_total\_centred"] and df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$stimulus == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 554, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -1.586608 1.586608  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862855 -0.000000000000002862855

# p > .05 - sticsa is independent of stimulus  
  
# sticsa and time  
t\_test\_independence\_sticsa\_time\_ext\_sacc\_amplitude <-   
 t.test(  
 df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$time == "1", "sticsa\_total\_centred"],  
 df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$time == "-1", "sticsa\_total\_centred"],   
 var.equal = TRUE  
 )  
t\_test\_independence\_sticsa\_time\_ext\_sacc\_amplitude

##   
## Two Sample t-test  
##   
## data: df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$time == "1", "sticsa\_total\_centred"] and df\_long\_ext\_sacc\_amplitude[df\_long\_ext\_sacc\_amplitude$time == "-1", "sticsa\_total\_centred"]  
## t = 0, df = 554, p-value = 1  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## -1.586608 1.586608  
## sample estimates:  
## mean of x mean of y   
## -0.000000000000002862855 -0.000000000000002862855

# p > .05 - sticsa is independent of time

## *Homogeneity of Regression Slopes*

####### check homogeneity of regression slopes  
  
###### fixation count  
### acquisition  
homogeneity\_regression\_slopes\_acq\_fix\_count <-  
df\_long\_acq\_fix\_count %>%  
 anova\_test(fix\_count ~ sticsa\_total\_centred + iu\_group + stimulus + iu\_group\*stimulus +   
 sticsa\_total\_centred\*iu\_group + sticsa\_total\_centred\*stimulus +  
 sticsa\_total\_centred\*iu\_group\*stimulus)

## Coefficient covariances computed by hccm()

homogeneity\_regression\_slopes\_acq\_fix\_count

## ANOVA Table (type II tests)  
##   
## Effect DFn DFd F p p<.05 ges  
## 1 sticsa\_total\_centred 1 270 0.114 0.736 0.0004210  
## 2 iu\_group 1 270 6.146 0.014 \* 0.0220000  
## 3 stimulus 1 270 0.957 0.329 0.0040000  
## 4 iu\_group:stimulus 1 270 0.103 0.749 0.0003810  
## 5 sticsa\_total\_centred:iu\_group 1 270 3.336 0.069 0.0120000  
## 6 sticsa\_total\_centred:stimulus 1 270 0.154 0.695 0.0005710  
## 7 sticsa\_total\_centred:iu\_group:stimulus 1 270 0.021 0.885 0.0000783

# p-values > .05: no interactions between STICSA and grouping variables   
  
### extinction  
homogeneity\_regression\_slopes\_ext\_fix\_count <-  
df\_long\_ext\_fix\_count %>%  
 anova\_test(fix\_count ~ sticsa\_total\_centred + iu\_group + stimulus + time + iu\_group\*stimulus +   
 iu\_group\*time + stimulus\*time + sticsa\_total\_centred\*iu\_group +   
 sticsa\_total\_centred\*stimulus + sticsa\_total\_centred\*time +  
 sticsa\_total\_centred\*iu\_group\*stimulus + sticsa\_total\_centred\*iu\_group\*stimulus\*time)

## Coefficient covariances computed by hccm()

homogeneity\_regression\_slopes\_ext\_fix\_count

## ANOVA Table (type II tests)  
##   
## Effect DFn DFd F p p<.05  
## 1 sticsa\_total\_centred 1 540 1.391000 0.239000   
## 2 iu\_group 1 540 14.015000 0.000201 \*  
## 3 stimulus 1 540 0.866000 0.353000   
## 4 time 1 540 1.996000 0.158000   
## 5 iu\_group:stimulus 1 540 0.988000 0.321000   
## 6 iu\_group:time 1 540 1.272000 0.260000   
## 7 stimulus:time 1 540 0.013000 0.910000   
## 8 sticsa\_total\_centred:iu\_group 1 540 0.719000 0.397000   
## 9 sticsa\_total\_centred:stimulus 1 540 0.238000 0.626000   
## 10 sticsa\_total\_centred:time 1 540 0.000156 0.990000   
## 11 sticsa\_total\_centred:iu\_group:stimulus 1 540 0.024000 0.876000   
## 12 sticsa\_total\_centred:iu\_group:time 1 540 0.335000 0.563000   
## 13 sticsa\_total\_centred:stimulus:time 1 540 0.166000 0.683000   
## 14 iu\_group:stimulus:time 1 540 0.008000 0.928000   
## 15 sticsa\_total\_centred:iu\_group:stimulus:time 1 540 0.103000 0.748000   
## ges  
## 1 0.00300000  
## 2 0.02500000  
## 3 0.00200000  
## 4 0.00400000  
## 5 0.00200000  
## 6 0.00200000  
## 7 0.00002370  
## 8 0.00100000  
## 9 0.00044000  
## 10 0.00000029  
## 11 0.00004500  
## 12 0.00062000  
## 13 0.00030800  
## 14 0.00001510  
## 15 0.00019100

# p-values > .05: no interactions between STICSA and grouping variables   
  
  
###### fixation duration  
### acquisition  
homogeneity\_regression\_slopes\_acq\_fix\_duration\_log <-  
df\_long\_acq\_fix\_duration\_log %>%  
 anova\_test(fix\_duration\_log ~ sticsa\_total\_centred + iu\_group + stimulus + iu\_group\*stimulus +   
 sticsa\_total\_centred\*iu\_group + sticsa\_total\_centred\*stimulus +  
 sticsa\_total\_centred\*iu\_group\*stimulus)

## Coefficient covariances computed by hccm()

homogeneity\_regression\_slopes\_acq\_fix\_duration\_log

## ANOVA Table (type II tests)  
##   
## Effect DFn DFd F p p<.05  
## 1 sticsa\_total\_centred 1 270 0.515000 0.473   
## 2 iu\_group 1 270 7.485000 0.007 \*  
## 3 stimulus 1 270 0.207000 0.650   
## 4 iu\_group:stimulus 1 270 0.123000 0.727   
## 5 sticsa\_total\_centred:iu\_group 1 270 1.643000 0.201   
## 6 sticsa\_total\_centred:stimulus 1 270 0.030000 0.863   
## 7 sticsa\_total\_centred:iu\_group:stimulus 1 270 0.000261 0.987   
## ges  
## 1 0.002000000  
## 2 0.027000000  
## 3 0.000766000  
## 4 0.000454000  
## 5 0.006000000  
## 6 0.000111000  
## 7 0.000000967

# p-values > .05: no interactions between STICSA and grouping variables   
  
### extinction  
homogeneity\_regression\_slopes\_ext\_fix\_duration\_log <-  
df\_long\_ext\_fix\_duration\_log %>%  
 anova\_test(fix\_duration\_log ~ sticsa\_total\_centred + iu\_group + stimulus + time + iu\_group\*stimulus +   
 iu\_group\*time + stimulus\*time + sticsa\_total\_centred\*iu\_group +   
 sticsa\_total\_centred\*stimulus + sticsa\_total\_centred\*time +  
 sticsa\_total\_centred\*iu\_group\*stimulus + sticsa\_total\_centred\*iu\_group\*stimulus\*time)

## Coefficient covariances computed by hccm()

homogeneity\_regression\_slopes\_ext\_fix\_duration\_log

## ANOVA Table (type II tests)  
##   
## Effect DFn DFd F p  
## 1 sticsa\_total\_centred 1 540 0.004000 0.951000000  
## 2 iu\_group 1 540 26.131000 0.000000444  
## 3 stimulus 1 540 0.121000 0.728000000  
## 4 time 1 540 1.651000 0.199000000  
## 5 iu\_group:stimulus 1 540 1.492000 0.222000000  
## 6 iu\_group:time 1 540 0.193000 0.661000000  
## 7 stimulus:time 1 540 0.035000 0.852000000  
## 8 sticsa\_total\_centred:iu\_group 1 540 0.228000 0.633000000  
## 9 sticsa\_total\_centred:stimulus 1 540 0.054000 0.816000000  
## 10 sticsa\_total\_centred:time 1 540 0.127000 0.722000000  
## 11 sticsa\_total\_centred:iu\_group:stimulus 1 540 0.000829 0.977000000  
## 12 sticsa\_total\_centred:iu\_group:time 1 540 0.189000 0.664000000  
## 13 sticsa\_total\_centred:stimulus:time 1 540 0.041000 0.839000000  
## 14 iu\_group:stimulus:time 1 540 0.070000 0.791000000  
## 15 sticsa\_total\_centred:iu\_group:stimulus:time 1 540 0.010000 0.919000000  
## p<.05 ges  
## 1 0.00000709  
## 2 \* 0.04600000  
## 3 0.00022500  
## 4 0.00300000  
## 5 0.00300000  
## 6 0.00035700  
## 7 0.00006460  
## 8 0.00042200  
## 9 0.00010100  
## 10 0.00023400  
## 11 0.00000153  
## 12 0.00035000  
## 13 0.00007630  
## 14 0.00013100  
## 15 0.00001900

# p-values > .05: no interactions between STICSA and grouping variables   
  
  
###### saccade amplitude  
### acquisition  
homogeneity\_regression\_slopes\_acq\_sacc\_amplitude <-  
df\_long\_acq\_sacc\_amplitude %>%  
 anova\_test(sacc\_amplitude ~ sticsa\_total\_centred + iu\_group + stimulus + iu\_group\*stimulus +   
 sticsa\_total\_centred\*iu\_group + sticsa\_total\_centred\*stimulus +  
 sticsa\_total\_centred\*iu\_group\*stimulus)

## Warning: NA detected in rows: 234,259.  
## Removing this rows before the analysis.

## Coefficient covariances computed by hccm()

homogeneity\_regression\_slopes\_acq\_sacc\_amplitude

## ANOVA Table (type II tests)  
##   
## Effect DFn DFd F p p<.05 ges  
## 1 sticsa\_total\_centred 1 268 0.018 0.894 0.0000664  
## 2 iu\_group 1 268 3.272 0.072 0.0120000  
## 3 stimulus 1 268 0.298 0.585 0.0010000  
## 4 iu\_group:stimulus 1 268 0.162 0.688 0.0006040  
## 5 sticsa\_total\_centred:iu\_group 1 268 0.038 0.846 0.0001410  
## 6 sticsa\_total\_centred:stimulus 1 268 0.166 0.684 0.0006180  
## 7 sticsa\_total\_centred:iu\_group:stimulus 1 268 0.136 0.713 0.0005060

# p-values > .05: no interactions between STICSA and grouping variables   
  
### extinction  
homogeneity\_regression\_slopes\_ext\_sacc\_amplitude <-  
df\_long\_ext\_sacc\_amplitude %>%  
 anova\_test(sacc\_amplitude ~ sticsa\_total\_centred + iu\_group + stimulus + time + iu\_group\*stimulus +   
 iu\_group\*time + stimulus\*time + sticsa\_total\_centred\*iu\_group +   
 sticsa\_total\_centred\*stimulus + sticsa\_total\_centred\*time +  
 sticsa\_total\_centred\*iu\_group\*stimulus + sticsa\_total\_centred\*iu\_group\*stimulus\*time)

## Warning: NA detected in rows: 116,181,301.  
## Removing this rows before the analysis.

## Coefficient covariances computed by hccm()

homogeneity\_regression\_slopes\_ext\_sacc\_amplitude

## ANOVA Table (type II tests)  
##   
## Effect DFn DFd F p p<.05  
## 1 sticsa\_total\_centred 1 537 2.227 0.136   
## 2 iu\_group 1 537 3.433 0.064   
## 3 stimulus 1 537 0.267 0.605   
## 4 time 1 537 0.125 0.724   
## 5 iu\_group:stimulus 1 537 0.682 0.409   
## 6 iu\_group:time 1 537 0.163 0.686   
## 7 stimulus:time 1 537 0.033 0.855   
## 8 sticsa\_total\_centred:iu\_group 1 537 7.992 0.005 \*  
## 9 sticsa\_total\_centred:stimulus 1 537 0.097 0.755   
## 10 sticsa\_total\_centred:time 1 537 0.420 0.517   
## 11 sticsa\_total\_centred:iu\_group:stimulus 1 537 1.339 0.248   
## 12 sticsa\_total\_centred:iu\_group:time 1 537 0.209 0.648   
## 13 sticsa\_total\_centred:stimulus:time 1 537 0.202 0.653   
## 14 iu\_group:stimulus:time 1 537 0.407 0.524   
## 15 sticsa\_total\_centred:iu\_group:stimulus:time 1 537 1.359 0.244   
## ges  
## 1 0.0040000  
## 2 0.0060000  
## 3 0.0004970  
## 4 0.0002330  
## 5 0.0010000  
## 6 0.0003040  
## 7 0.0000619  
## 8 0.0150000  
## 9 0.0001810  
## 10 0.0007810  
## 11 0.0020000  
## 12 0.0003890  
## 13 0.0003760  
## 14 0.0007570  
## 15 0.0030000

# p-values > .05: no interactions between STICSA and grouping variables, except for  
# sticsa\*iu p = .005

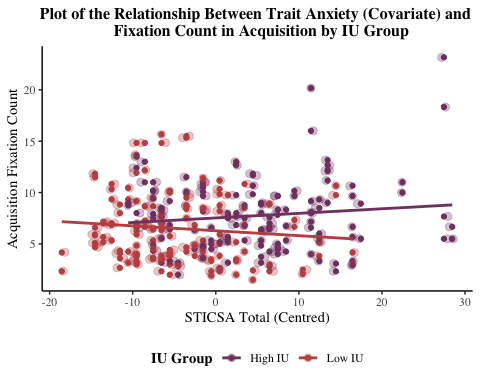
## *Linearity Between Covariate and Outcome Variables*

### *Fixation Count*

#### *Acquisition*

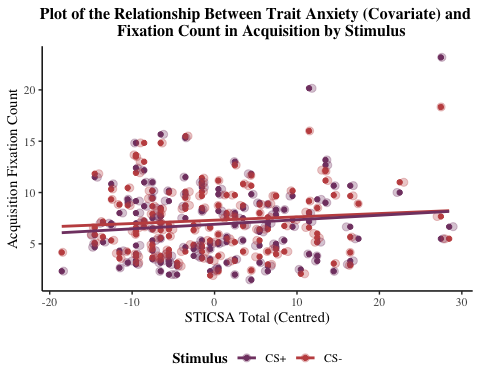
## this is at each level of grouping variable.   
# check by computing grouped scatterplot of covariate and outcome variable   
  
# sticsa and IU group  
scatterplot\_acq\_fix\_count\_sticsa\_centred\_by\_iu <-   
 ggplot(df\_long\_acq\_fix\_count,aes(x = sticsa\_total\_centred, y = fix\_count,   
 colour = iu\_group)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Count in Acquisition by IU Group",  
 x = "STICSA Total (Centred)",   
 y = "Acquisition Fixation Count") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(colour = "IU Group") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_acq\_fix\_count\_sticsa\_centred\_by\_iu)

## `geom\_smooth()` using formula 'y ~ x'



# relationship between STICSA and fixation count appears linear at both levels of IU  
  
# sticsa and stimulus  
scatterplot\_acq\_fix\_count\_sticsa\_centred\_by\_stimulus <-   
 ggplot(df\_long\_acq\_fix\_count,aes(x = sticsa\_total\_centred, y = fix\_count,   
 colour = stimulus)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Count in Acquisition by Stimulus",  
 x = "STICSA Total (Centred)",   
 y = "Acquisition Fixation Count") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("CS-", "CS+")) +  
 labs(colour = "Stimulus") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_acq\_fix\_count\_sticsa\_centred\_by\_stimulus)

## `geom\_smooth()` using formula 'y ~ x'

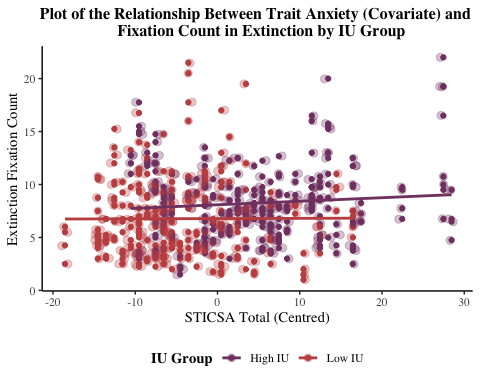


# relationship between STICSA and fixation count appears linear at both levels of stimulus

#### *Extinction*

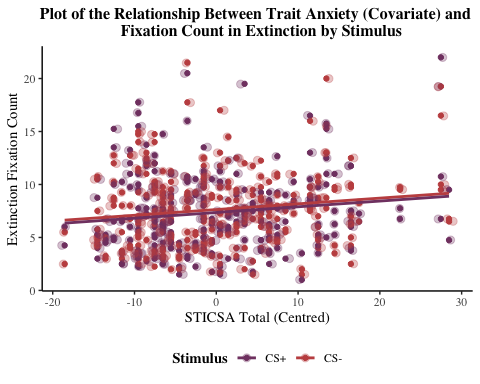
# sticsa and IU group  
scatterplot\_ext\_fix\_count\_sticsa\_centred\_by\_iu <-   
 ggplot(df\_long\_ext\_fix\_count,aes(x = sticsa\_total\_centred, y = fix\_count,   
 colour = iu\_group)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Count in Extinction by IU Group",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Fixation Count") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(colour = "IU Group") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_fix\_count\_sticsa\_centred\_by\_iu)

## `geom\_smooth()` using formula 'y ~ x'



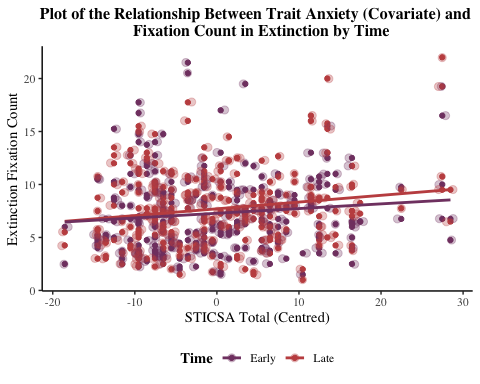
# relationship between STICSA and fixation count appears linear at both levels of IU  
  
# sticsa and stimulus  
scatterplot\_ext\_fix\_count\_sticsa\_centred\_by\_stimulus <-   
 ggplot(df\_long\_ext\_fix\_count,aes(x = sticsa\_total\_centred, y = fix\_count,   
 colour = stimulus)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Count in Extinction by Stimulus",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Fixation Count") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("CS-", "CS+")) +  
 labs(colour = "Stimulus") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_fix\_count\_sticsa\_centred\_by\_stimulus)

## `geom\_smooth()` using formula 'y ~ x'



# relationship between STICSA and fixation count appears linear at both levels of stimulus  
  
# sticsa and time  
scatterplot\_ext\_fix\_count\_sticsa\_centred\_by\_time <-   
 ggplot(df\_long\_ext\_fix\_count,aes(x = sticsa\_total\_centred, y = fix\_count,   
 colour = time)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Count in Extinction by Time",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Fixation Count") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Late", "Early")) +  
 labs(colour = "Time") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_fix\_count\_sticsa\_centred\_by\_time)

## `geom\_smooth()` using formula 'y ~ x'



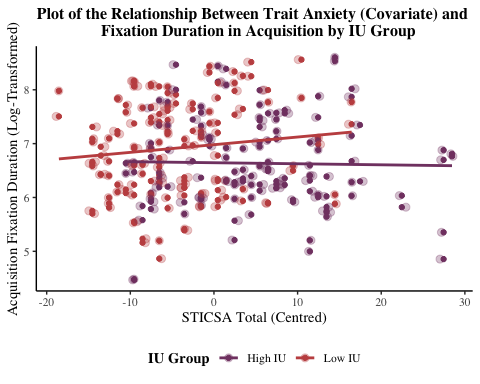
# relationship between STICSA and fixation count appears linear at both levels of time

### *Fixation Duration - Log Transformed*

#### *Acquisition*

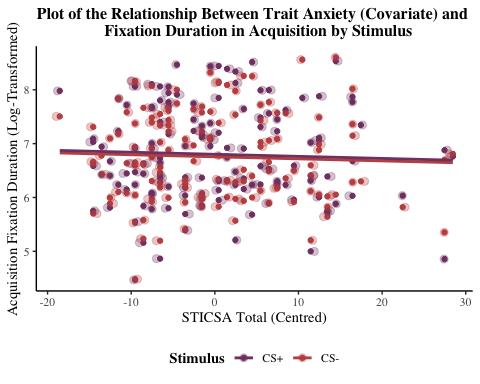
# sticsa and IU group  
scatterplot\_acq\_fix\_duration\_log\_sticsa\_centred\_by\_iu <-   
 ggplot(df\_long\_acq\_fix\_duration\_log, aes(x = sticsa\_total\_centred, y = fix\_duration\_log,   
 colour = iu\_group)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Duration in Acquisition by IU Group",  
 x = "STICSA Total (Centred)",   
 y = "Acquisition Fixation Duration (Log-Transformed)") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(colour = "IU Group") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_acq\_fix\_duration\_log\_sticsa\_centred\_by\_iu)

## `geom\_smooth()` using formula 'y ~ x'



# relationship between STICSA and fixation count appears linear at both levels of IU  
  
# sticsa and stimulus  
scatterplot\_acq\_fix\_duration\_log\_sticsa\_centred\_by\_stimulus <-   
 ggplot(df\_long\_acq\_fix\_duration\_log, aes(x = sticsa\_total\_centred, y = fix\_duration\_log,   
 colour = stimulus)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Duration in Acquisition by Stimulus",  
 x = "STICSA Total (Centred)",   
 y = "Acquisition Fixation Duration (Log-Transformed)") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("CS-", "CS+")) +  
 labs(colour = "Stimulus") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_acq\_fix\_duration\_log\_sticsa\_centred\_by\_stimulus)

## `geom\_smooth()` using formula 'y ~ x'

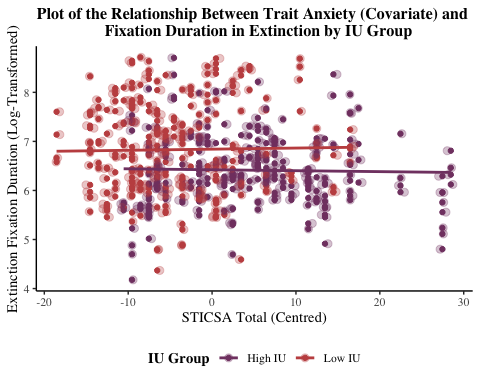


# relationship between STICSA and fixation count appears linear at both levels of stimulus

#### *Extinction*

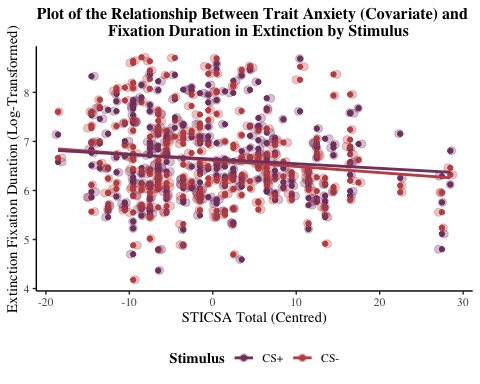
# sticsa and IU group  
scatterplot\_ext\_fix\_duration\_log\_sticsa\_centred\_by\_iu <-   
 ggplot(df\_long\_ext\_fix\_duration\_log, aes(x = sticsa\_total\_centred, y = fix\_duration\_log,   
 colour = iu\_group)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Duration in Extinction by IU Group",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Fixation Duration (Log-Transformed)") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(colour = "IU Group") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_fix\_duration\_log\_sticsa\_centred\_by\_iu)

## `geom\_smooth()` using formula 'y ~ x'



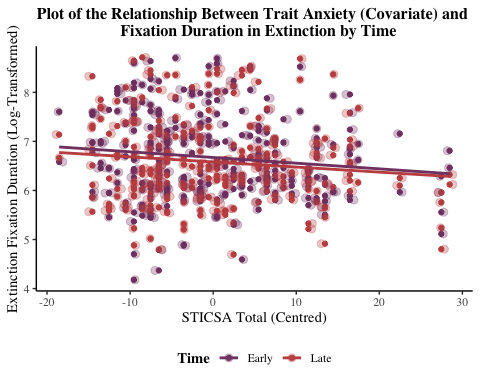
# relationship between STICSA and fixation count appears linear at both levels of IU  
  
# sticsa and stimulus  
scatterplot\_ext\_fix\_duration\_log\_sticsa\_centred\_by\_stimulus <-   
 ggplot(df\_long\_ext\_fix\_duration\_log, aes(x = sticsa\_total\_centred, y = fix\_duration\_log,   
 colour = stimulus)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Duration in Extinction by Stimulus",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Fixation Duration (Log-Transformed)") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("CS-", "CS+")) +  
 labs(colour = "Stimulus") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_fix\_duration\_log\_sticsa\_centred\_by\_stimulus)

## `geom\_smooth()` using formula 'y ~ x'



# relationship between STICSA and fixation count appears linear at both levels of stimulus  
  
# sticsa and time  
scatterplot\_ext\_fix\_duration\_log\_sticsa\_centred\_by\_time <-   
 ggplot(df\_long\_ext\_fix\_duration\_log, aes(x = sticsa\_total\_centred, y = fix\_duration\_log,   
 colour = time)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Fixation Duration in Extinction by Time",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Fixation Duration (Log-Transformed)") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Late", "Early")) +  
 labs(colour = "Time") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_fix\_duration\_log\_sticsa\_centred\_by\_time)

## `geom\_smooth()` using formula 'y ~ x'



# relationship between STICSA and fixation count appears linear at both levels of time

### *Saccade Amplitude*

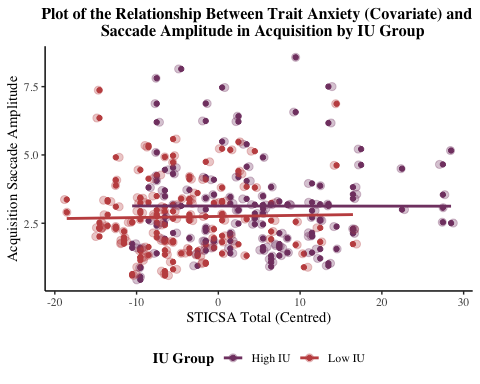
#### *Acquisition*

# sticsa and IU group  
scatterplot\_acq\_sacc\_amplitude\_sticsa\_centred\_by\_iu <-   
 ggplot(df\_long\_acq\_sacc\_amplitude, aes(x = sticsa\_total\_centred, y = sacc\_amplitude,   
 colour = iu\_group)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Saccade Amplitude in Acquisition by IU Group",  
 x = "STICSA Total (Centred)",   
 y = "Acquisition Saccade Amplitude") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(colour = "IU Group") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_acq\_sacc\_amplitude\_sticsa\_centred\_by\_iu)

## `geom\_smooth()` using formula 'y ~ x'

## Warning: Removed 2 rows containing non-finite values (stat\_smooth).

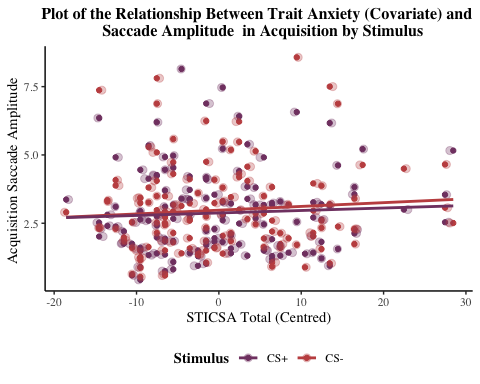
## Warning: Removed 2 rows containing missing values (geom\_point).  
  
## Warning: Removed 2 rows containing missing values (geom\_point).



# relationship between STICSA and fixation count appears linear at both levels of IU  
  
# sticsa and stimulus  
scatterplot\_acq\_sacc\_amplitude\_sticsa\_centred\_by\_stimulus <-   
 ggplot(df\_long\_acq\_sacc\_amplitude, aes(x = sticsa\_total\_centred, y = sacc\_amplitude,   
 colour = stimulus)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Saccade Amplitude in Acquisition by Stimulus",  
 x = "STICSA Total (Centred)",   
 y = "Acquisition Saccade Amplitude ") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("CS-", "CS+")) +  
 labs(colour = "Stimulus") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_acq\_sacc\_amplitude\_sticsa\_centred\_by\_stimulus)

## `geom\_smooth()` using formula 'y ~ x'

## Warning: Removed 2 rows containing non-finite values (stat\_smooth).  
  
## Warning: Removed 2 rows containing missing values (geom\_point).  
  
## Warning: Removed 2 rows containing missing values (geom\_point).



# relationship between STICSA and fixation count appears linear at both levels of stimulus

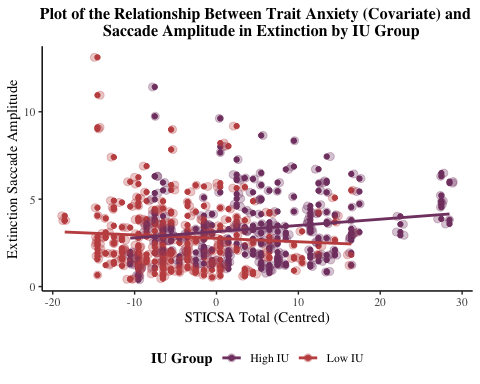
#### *Extinction*

# sticsa and IU group  
scatterplot\_ext\_sacc\_amplitude\_sticsa\_centred\_by\_iu <-   
 ggplot(df\_long\_ext\_sacc\_amplitude, aes(x = sticsa\_total\_centred, y = sacc\_amplitude,   
 colour = iu\_group)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Saccade Amplitude in Extinction by IU Group",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Saccade Amplitude") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Low IU", "High IU")) +  
 labs(colour = "IU Group") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_sacc\_amplitude\_sticsa\_centred\_by\_iu)

## `geom\_smooth()` using formula 'y ~ x'

## Warning: Removed 3 rows containing non-finite values (stat\_smooth).

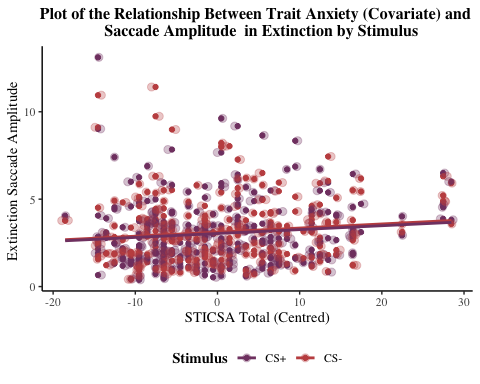
## Warning: Removed 3 rows containing missing values (geom\_point).  
  
## Warning: Removed 3 rows containing missing values (geom\_point).



# relationship between STICSA and fixation count appears linear at both levels of IU  
# there does appear to be an interaction (with high IU having higher  
# saccde amplitude as levels of trait anxiety increase, and low IU showing  
# opposite pattern)  
  
# sticsa and stimulus  
scatterplot\_ext\_sacc\_amplitude\_sticsa\_centred\_by\_stimulus <-   
 ggplot(df\_long\_ext\_sacc\_amplitude, aes(x = sticsa\_total\_centred, y = sacc\_amplitude,   
 colour = stimulus)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Saccade Amplitude in Extinction by Stimulus",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Saccade Amplitude ") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("CS-", "CS+")) +  
 labs(colour = "Stimulus") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_sacc\_amplitude\_sticsa\_centred\_by\_stimulus)

## `geom\_smooth()` using formula 'y ~ x'

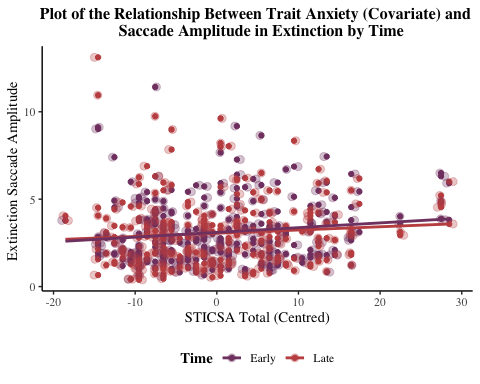
## Warning: Removed 3 rows containing non-finite values (stat\_smooth).  
  
## Warning: Removed 3 rows containing missing values (geom\_point).  
  
## Warning: Removed 3 rows containing missing values (geom\_point).



# relationship between STICSA and fixation count appears linear at both levels of stimulus  
  
# sticsa and time  
scatterplot\_ext\_sacc\_amplitude\_sticsa\_centred\_by\_time <-   
 ggplot(df\_long\_ext\_sacc\_amplitude, aes(x = sticsa\_total\_centred, y = sacc\_amplitude,   
 colour = time)) +  
 geom\_point() +  
 geom\_jitter(width = .5, alpha = .30, size = 2.5) +  
 geom\_smooth(method = lm, se = FALSE) +  
 labs(title = "Plot of the Relationship Between Trait Anxiety (Covariate) and   
 Saccade Amplitude in Extinction by Time",  
 x = "STICSA Total (Centred)",   
 y = "Extinction Saccade Amplitude ") +  
 theme\_classic() +  
 theme(plot.title = element\_text(face = "bold", hjust = 0.5, size = 12)) +  
 theme(text = element\_text(family = "serif")) +  
 guides(colour = guide\_legend(reverse = TRUE)) +  
 scale\_colour\_manual(values = c("#c45150", "#824372"), labels = c("Late", "Early")) +  
 labs(colour = "Time") +  
 theme(legend.position = "bottom", legend.title = element\_text(face = "bold"))   
  
print(scatterplot\_ext\_sacc\_amplitude\_sticsa\_centred\_by\_time)

## `geom\_smooth()` using formula 'y ~ x'

## Warning: Removed 3 rows containing non-finite values (stat\_smooth).  
  
## Warning: Removed 3 rows containing missing values (geom\_point).  
  
## Warning: Removed 3 rows containing missing values (geom\_point).



# relationship between STICSA and fixation count appears linear at both levels of time