

```

---
title: "Raw Data Merge"
output:
  html_document:
    df_print: paged
  html_notebook: default
  pdf_document: default
---

```

```

```{r}
library(tidyverse)
library(googleheets4)
library(qualtrics)
library(lubridate)
```

```

Note: the Qualtrics pretest data still has participant names, so this script cannot be fully executed without the Qualtrics credentials, which are stored elsewhere.

```

## Load list of valid IDs
```{r download}
gs4_auth(path=Sys.getenv("SHINY_PROTOCOL"))

#gs4_auth(email="russell.g.almond@gmail.com",
scopes="https://www.googleapis.com/auth/spreadsheets")
#gs4_deauth()

ValidIDs <- read_sheet("1lBp0LhNdSf0r1OtWC2cyqInB2npLbjH0KUMh4z_nzfM")$StudyID
AllIDs <- read_sheet("1BPkBMEE5DOtZ3MBt5C1ab4djddAOhUyAIPMiR4WRhBQ")

AllIDs %>% filter(StudyID %in% ValidIDs) -> PPIDs
AllIDs %>% filter(!(StudyID %in% ValidIDs)) -> InvIDs
```

# Survey Metadata

Experiment in trying to mine the QSF information to automagically build data
dictionary.

```{r QualtricsSurveyIDs}
qualtricsIDs <- c(PRE="SV_bkALP80IYWnEgkK",ECT="SV_3BKybDb3f2v4QU6",
  POT="SV_1MotNHuldPEAJnM",Assent="SV_07esYssbhDccXs1",
  Consent="SV_8vsIO1kl7DSPMG", PosttestA="SV_aeBB1Tw7bYKeIqq",
  PosttestB="SV_4ISgzDcDm6BLiom", PretestA="SV_4IpHb99rMIEKtIq",
  PretestB="SV_6eTzhLhhnhXrPmK")
```

## Question Titles
```{r grabMetadata}
pretestQ <- survey_questions(qualtricsIDs["PRE"])
potQ <- survey_questions(qualtricsIDs["POT"])
ectQ <- survey_questions(qualtricsIDs["ECT"])
posttestB <- survey_questions(qualtricsIDs["PosttestB"])
```

```{r cleanHTML}

```

```

pretestQuestions <- gsub("</?p>", "", pretestQ$question)
names(pretestQuestions) <- pretestQ$qname
potQuestions <- gsub("</?p>", "", potQ$question)
names(potQuestions) <- potQ$qname
ectQuestions <- gsub("</?p>", "", ectQ$question)
names(ectQuestions) <- ectQ$qname
postBQuestions <- gsub("</?p>", "", posttestB$question)
names(postBQuestions) <- posttestB$qname
```

```

## Qualtrics QSF

```

```{r LoadQSF}
preQSF <- jsonlite::fromJSON("PP_IES_Pretest.qsf", FALSE)
potQSF <- jsonlite::fromJSON("PP_IES_Posttest_POT (2).qsf", FALSE)
ectQSF <- jsonlite::fromJSON("PP_IES_Posttest_ECT.qsf", FALSE)
postBQSF <- jsonlite::fromJSON("Posttest_B_Summer_Camp.qsf", FALSE)
```

```

Fetch the elements marked "SQ". Use the DataExportTag as the name, this should allow searching the metadata by name.

```

```{r preSQ}
preSQ <- lapply(preQSF$SurveyElements[which(sapply(preQSF$SurveyElements,
  function(el) el$Element=="SQ"))],
               function(sq) sq$Payload)
names(preSQ) <- sapply(preSQ, function(q) q$DataExportTag)
potSQ <- lapply(potQSF$SurveyElements[which(sapply(potQSF$SurveyElements,
  function(el) el$Element=="SQ"))],
               function(sq) sq$Payload)
names(potSQ) <- sapply(potSQ, function(q) q$DataExportTag)
ectSQ <- lapply(ectQSF$SurveyElements[which(sapply(ectQSF$SurveyElements,
  function(el) el$Element=="SQ"))],
               function(sq) sq$Payload)
names(ectSQ) <- sapply(ectSQ, function(q) q$DataExportTag)
postBSQ <- lapply(postBQSF$SurveyElements[which(sapply(postBQSF$SurveyElements,
   function(el) el$Element=="SQ"))],
                 function(sq) sq$Payload)
names(postBSQ) <- sapply(postBSQ, function(q) q$DataExportTag)
```

```

## Names for IMI questions

```

```{r IMI}
choices <- ectSQ$IMI$Choices
IMI.choices <- rep(NA_character_, max(as.numeric(names(choices))))
for (choi in names(choices)) {
  IMI.choices[as.numeric(choi)] <- choices[[choi]]$Display
}
chorder <- as.numeric(unlist(ectSQ$IMI$ChoiceOrder))
IMI.stems <- IMI.choices[chorder]
names(IMI.stems) <- paste("IMI", 1L:length(IMI.stems), sep="_")
IMI.stems <- gsub("\t", "", IMI.stems) # Clean extra tabs.
```

```

## Names for PA questions.

PA questions only appear on ECD form.

```
```{r PA.metadata}
PA.names <- paste("PA",1:7,sep="")
PA.stems <- sapply(ectSQ[PA.names],function (sq) sq$QuestionText)
PA.labels <- sapply(ectSQ[PA.names],function(sq) unlist(sq$Choices))
PA.labels
```
```

## Names for the ethnicity questions

```
```{r ethnicityMetadata}
ethnames <- preSQ$Ethnicity$Choices
ethcolnames <- paste("Ethnicity",names(ethnames),sep="_")
ethnames <- unlist(ethnames)
ethnames <- ethnames[grepl("Display",names(ethnames))]
names(ethnames) <- ethcolnames
ethnames
```
```

## Extract the keys

```
```{r getKeys}
preKey <- sapply(preSQ,function(sq) {
  if (is.null(sq$GradingData) || length(sq$GradingData)==0L) return (NA)
  as.numeric(sq$GradingData[[1]]$ChoiceID)
})
preKey <- preKey[!is.na(preKey)]
potKey <- sapply(potSQ,function(sq) {
  if (is.null(sq$GradingData) || length(sq$GradingData)==0L) return (NA)
  as.numeric(sq$GradingData[[1]]$ChoiceID)
})
potKey <- potKey[!is.na(potKey)]
ectKey <- sapply(ectSQ,function(sq) {
  if (is.null(sq$GradingData) || length(sq$GradingData)==0L) return (NA)
  as.numeric(sq$GradingData[[1]]$ChoiceID)
})
ectKey <- ectKey[!is.na(ectKey)]
postBKey <- sapply(postBSQ,function(sq) {
  if (is.null(sq$GradingData) || length(sq$GradingData)==0L) return (NA)
  as.numeric(sq$GradingData[[1]]$ChoiceID)
})
postBKey <- postBKey[!is.na(postBKey)]
```
```

```
```{r KeyAndFormCode}
keyID <- "1fagyGKc30Fx2ORwcWoULJ_dYZWqwPwJWCc-xo1pfV2o"
physics.key <- read_sheet(keyID,"Physics")
IMI.scales <- read_sheet(keyID,"IMI")
PA.rev <- read_sheet(keyID,"PA")
```
```

I have key information from two sources, Qualtrics and external file. Check to see if they are the same.

```
```{r keyCheck}
formA.keys <- filter(physics.key,Form=="A") %>% select(ID,Key)
```

```

formA.keys$prekey <- preKey[formA.keys$ID]
formA.keys$potKey <- potKey[formA.keys$ID]
formA.keys$ectKey <- ectKey[formA.keys$ID]
formA.keys
```

```

No key mismatches, but column labeling issue. For some reason, `A-FQ10` is called `AFQ10` in pretest.

```

```{r getChoices}
choicesToFactor <- function (sq, col) {
  levels <- as.numeric(names(sq$Choices))
  labels <- sapply(sq$Choices, function(c) c$Display)
  factor(col,levels,labels)
}
```

```

# Load The Qualtrics Data

Exclude metadata fields, fields containing PID and "DO" fields which give presentation ordering.

```

```{r LoadPrePost}
pretest <- fetch_survey(qualtricsIDs["PRE"],force=TRUE,
                        label=FALSE,convert=FALSE) %>%
  filter(ID %in% ValidIDs) %>%
  select(!(StartDate:IPAddress)) %>%
  select(!(ResponseId:UserLanguage)) %>%
  select(!(First:Q80)) %>%
  select(!contains("DO"))
names(pretest)[5] <- "StudyID"
pretest <- filter(pretest,!is.na(StudyID))

ecttest <- fetch_survey(qualtricsIDs["ECT"],force=TRUE,
                        label=FALSE, convert=FALSE) %>%
  filter(UserID1 %in% ValidIDs) %>%
  select(!(StartDate:IPAddress)) %>%
  select(!(ResponseId:UserLanguage)) %>%
  select(!contains("DO"))
names(ecttest)[5] <- "StudyID"
ecttest <- filter(ecttest,!is.na(StudyID))

pottest <- fetch_survey(qualtricsIDs["POT"], force=TRUE,
                        label=FALSE, convert=FALSE) %>%
  filter(UserID1 %in% ValidIDs) %>%
  select(!(StartDate:IPAddress)) %>%
  select(!(ResponseId:UserLanguage)) %>%
  select(!contains("DO"))
names(pottest)[5] <- "StudyID"
pottest <- filter(pottest,!is.na(StudyID))
```

```

## Clean duplicates.

```

```{r dupesbytest}
dupes.pre <- pretest$StudyID[duplicated(pretest$StudyID)]
"Pretest:"
dupes.pre
"ECT:"
dupes.ect <- ecttest$StudyID[duplicated(ecttest$StudyID)]
dupes.ect
"POT:"
dupes.pot <- pottest$StudyID[duplicated(pottest$StudyID)]
dupes.pot
```

```

### POT

```

```{r POTdupe}
pottest %>% filter(StudyID%in%dupes.pot)
```

```{r cleanPOT}
pottest <- filter(pottest,! (StudyID %in% dupes.pot) | Finished)
dupes.pot1 <- pottest$StudyID[duplicated(pottest$StudyID)]
pottest %>% filter(StudyID%in%dupes.pot1)
```

```

For A2079 and D3654 take the row with non-missing PA1  
 For F1562 Flip a coin.

```

```{r dropDups}
pottest <- filter(pottest,! (StudyID %in% c("A2079","D3654"))) | !is.na(PA1))
dupes.pot2 <- pottest$StudyID[duplicated(pottest$StudyID)]
for (dupID in dupes.pot2) {
  dupindex <- which(pottest$StudyID==dupID)
  dropindex <- sample(dupindex,length(dupindex)-1,replace=FALSE)
  pottest <- pottest[-dropindex,]
}
anyDuplicated(pottest$StudyID)
```

```

### ECT

```

```{r dupECT}

filter(ecttest, StudyID %in% dupes.ect) %>% arrange(StudyID)
```

Select the ones which are finished.
```{r cleanECT}
ecttest <- filter(ecttest,! (StudyID %in% dupes.ect) | Finished)
ecttest$StudyID[duplicated(ecttest$StudyID)]
```

```

### PRE

```

```{r dupPRE}

filter(pretest, StudyID %in% dupes.pre) %>%

```

```

  arrange(StudyID) %>% select(Progress, Finished, StudyID)
  ```
  ```{r cleanPRE}
  pretest <- filter(pretest,! (StudyID %in% dupes.pre) | Finished)
  dupes.pre1 <- pretest$StudyID[duplicated(pretest$StudyID)]
  filter(pretest,StudyID %in% dupes.pre1)
  ```
  Still one ambiguous case. Take the one with the longest duration.

  ```{r preDuration}
  for (dupID in dupes.pre1) {
    dupindex <- which(pretest$StudyID==dupID)
    dur <- pretest[dupindex,"Duration (in seconds)",drop=TRUE]
    dropindex <- dupindex[dur < max(dur)]
    pretest <- pretest[-dropindex,]
  }
  anyDuplicated(pretest$StudyID)
  ```

  ## Covert to factors and add labels.

  ### Demographics

  Only a few students checked Nonbinary or Other, so put them together. Also, call
  prefer not to say as NA.

  A couple of students put XX years old, instead of their age, so fix.

  ```{r demofix}
  pretest$Sex <- choicesToFactor(preSQ$Sex,pretest$Sex)
  pretest$Gender <- case_match(pretest$Sex,
                                "Male"~"Male",
                                "Female"~"Female",
                                c("Other","Nonbinary") ~ "Other",
                                .default=NA) %>%
    factor(levels=c("Male","Female","Other"))
  age <- pretest$Age
  age[!is.na(age)] <- regmatches(age[!is.na(age)],
                                regexpr("[[:digit:]]*",
                                age[!is.na(age)]))
  pretest$Age <- as.numeric(age)
  pretest$Gaming <- choicesToFactor(preSQ$Gaming,pretest$Gaming)
  pretest$Physics <- choicesToFactor(preSQ$Physics,pretest$Physics)
  ```

  Ethnicity. Collapse a couple of categories.
  ```{r ethnicityOther}
  pretest$Ethnicity_9_TEXT[!is.na(pretest$Ethnicity_9)]
  ```

  Qualtrics "helpfully" codes this value as 1/NA. Fix to true logicals.

  ```{r Logical}
  ethids <- grep("Ethnicity_",names(pretest))
  for (eth in ethids) {
    name <- names(pretest)[eth]

```

```

if (any(grepl("TEXT",name))) {
  names(pretest)[eth] <- "Other_TEXT"
} else {
  names(pretest)[eth] <- ethnames[name]
  pretest[[eth]] <- !is.na(pretest[[eth,drop=TRUE]])
}
}
pretest[,ethids]
```

```{r collapseEthnicity}
ethids1 <- ethids[-length(ethids)] ## Remove Other_TEXT
ethcols <- as.matrix(as.data.frame(pretest[,ethids1]))
ethnicity <- sapply(1L:nrow(pretest), function(irow) {
  paste(ethnames[ethcols[irow,]], collapse=",")
})
unique(ethnicity)
```

```

To get a single factor, collapse any combination into "mixed"

```

```{r recodeEth}
ethnicity[grepl(",",ethnicity)] <- "Mixed"
is.na(ethnicity) <- pretest$`Prefer not to say`
ethnicity <- as.factor(ethnicity)
summary(ethnicity)
pretest$Ethnicity <- ethnicity
```

```

### ### Physics Questions

Need to fix naming issue with Column A-FQ10. Also A-NQ1 and A-FQ4 metadata missing.

```

```{r Pretest Glitches}
names(pretest)[names(pretest)=="AFQ10"] <- "A-FQ10"
names(preKey)[names(preKey)=="AFQ10"] <- "A-FQ10"
names(preSQ)[names(preSQ)=="AFQ10"] <- "A-FQ10"
preSQ["A-NQ1"] <- potSQ["A-NQ1"]
preSQ["A-FQ4"] <- potSQ["A-FQ4"]
```

```{r encodeAndScore}
for (q in grep("A-",names(pretest),value=TRUE)) {
  vals <- pretest[[q]]
  pretest[[paste(q,"scored",sep="_")]] <- as.numeric(vals==preKey[q])
  pretest[[q]] <- choicesToFactor(preSQ[[q]],vals)
}
for (q in grep("A-",names(ecttest),value=TRUE)) {
  vals <- ecttest[[q]]
  ecttest[[paste(q,"scored",sep="_")]] <- as.numeric(vals==ectKey[q])
  ecttest[[q]] <- choicesToFactor(preSQ[[q]],vals)
}
for (q in grep("A-",names(pottest),value=TRUE)) {
  vals <- pottest[[q]]
  pottest[[paste(q,"scored",sep="_")]] <- as.numeric(vals==preKey[q])
}

```

```

    pottest[[q]] <- choicesToFactor(preSQ[[q]],vals)
  }
pretest
```

```{r subscales}
NearECTcols <- paste(physics.key$ID[physics.key$Form=="A" &
                        physics.key$`Near/Far`=="Near" &
                        physics.key$`HL Concept`=="EcT"],
                    "scored",sep="_")
FarECTcols <- paste(physics.key$ID[physics.key$Form=="A" &
                        physics.key$`Near/Far`=="Far" &
                        physics.key$`HL Concept`=="EcT"],
                    "scored",sep="_")
NearPOTcols <- paste(physics.key$ID[physics.key$Form=="A" &
                        physics.key$`Near/Far`=="Near" &
                        physics.key$`HL Concept`=="PoT"],
                    "scored",sep="_")
FarPOTcols <- paste(physics.key$ID[physics.key$Form=="A" &
                        physics.key$`Near/Far`=="Far" &
                        physics.key$`HL Concept`=="PoT"],
                    "scored",sep="_")
```

```{r subscores}
pretest %>%
  mutate(NearECT=rowSums(pretest[,NearECTcols],na.rm=TRUE),
         FarECT=rowSums(pretest[,FarECTcols],na.rm=TRUE),
         NearPOT=rowSums(pretest[,NearPOTcols],na.rm=TRUE),
         FarPOT=rowSums(pretest[,FarPOTcols],na.rm=TRUE)) %>%
  mutate(Near=NearECT+NearPOT,Far=FarECT+FarPOT,
         ECT=NearECT+FarECT, POT=NearPOT+FarPOT,
         PhysicsScore=NearECT+FarECT+NearPOT+FarPOT) ->
pretest
```

```{r subscoresPOST}
pottest %>%
  mutate(NearPOTpost=rowSums(pretest[,NearPOTcols],na.rm=TRUE),
         FarPOTpost=rowSums(pretest[,FarPOTcols],na.rm=TRUE)) %>%
  mutate(POTpost = NearPOTpost+FarPOTpost) ->
pottest
ecttest %>%
  mutate(NearECTpost=rowSums(ecttest[,NearECTcols],na.rm=TRUE),
         FarECTpost=rowSums(ecttest[,FarECTcols],na.rm=TRUE)) %>%
  mutate(ECTpost = NearECTpost+FarECTpost) ->
ecttest
```

### IMI Questions

IMI questions are only in the posttest.

```{r revcode}
for (col in IMI.scales$ID[IMI.scales$Reverse]) {
  pottest[[col]] <- 8-pottest[[col]]
  ecttest[[col]] <- 8-ecttest[[col]]
}

```



```

}
```

```{r IMISubScales}
pottest$IMI <- rowSums(pottest[,unique(IMI.scales$ID)],na.rm=TRUE)
ecttest$IMI <- rowSums(ecttest[,unique(IMI.scales$ID)],na.rm=TRUE)

for (scale in unique(IMI.scales$Scale)) {
  pottest[[paste("IMI",scale,sep="_")]] <-
    rowSums(pottest[,IMI.scales$ID[IMI.scales$Scale==scale]],na.rm=TRUE)
  ecttest[[paste("IMI",scale,sep="_")]] <-
    rowSums(ecttest[,IMI.scales$ID[IMI.scales$Scale==scale]],na.rm=TRUE)
}

```

```

### PA questions

Again, only in ECT and POT

UGH! Qualtrics was not consistent in the numeric values for these columns. Some were 1--5, some were 11--15, and one was 8--12. WTF. I went back and fixed the coding in Qualtrics.

```

```{r PArevcode}
for(paq in PA.rev$ID[PA.rev$Reversed]) {
  ecttest[[paq]] <- 5-ecttest[[paq]]
  pottest[[paq]] <- 5-pottest[[paq]]
}
ecttest$PA <- rowSums(ecttest[,PA.rev$ID],na.rm=TRUE)
pottest$PA <- rowSums(pottest[,PA.rev$ID],na.rm=TRUE)
pottest
ecttest
```

```

# Do the join

```

```{r join}
FSUSFall2022 <- select(PIDs,!Number) %>%
  full_join(pretest,by="StudyID",suffix=c("", ".pre")) %>%
  full_join(ecttest,by="StudyID",suffix=c("", ".ect")) %>%
  full_join(pottest,by="StudyID",suffix=c("", ".pot"))
```

```

## Check duplicates

```

```{r dupes}
dupes <- FSUSFall2022$StudyID[duplicated(FSUSFall2022$StudyID)]
dupes
```

```

Yay! Screening on Finished fixed the duplication problem.

# Output the data

```

```{r output}
outID <- "1DJf7Iidg-GvEXHkrfNPwGTB7VerLSTAUohzZYQOov9E"
write_sheet(FSUSFall2022,outID, sheet="Data")
write_sheet(data.frame(StudyID=dupes),outID,"Duplicate IDs")
write_csv(FSUSFall2022,"data/PPIESFall2022PrePost.csv")
```

## Generate cleaning script for the log data.

```{r invalid}
Xids <- c(InvIDs$ID1,InvIDs$ID2)
write_sheet(data.frame(Xids=Xids),outID,"DeleteThese")
```

This is now used to clean the log files.

# Merge data from Log Files

## BNScores
BNScores is keyed by uid, so we are ready to go. Kill the first two columns,
which are unneeded.

```{r BNScores}
BNScores <- read_csv("https://pluto.coe.fsu.edu/Proc4/dongle/data/stats-
BigStudy.csv")
BNScores <- select(BNScores,!any_of(c("...1","app")))
```

```{r writeBNScores}
write_sheet(BNScores, outID, "BNScores")
write_csv(BNScores,"data/PPIESFall2022BN.csv")
```

## Observables

observables has (uid,Context) as key, need to pivot wider. Timestamp isn't
really an
observable, but keep it anyway.

```{r cleanObs}
observables <-
read_csv("https://pluto.coe.fsu.edu/Proc4/dongle/data/ppObs.BigStudy.csv")
obsnames <- names(observables)[-(1:3)]
## Drop useless column and sort by timestamp
observables %>% select(!`...1`) %>% arrange(timestamp) -> observables
```

### Add Quits

```{r add quits}
observables$TrophyLevel[is.na(observables$TrophyLevel)] <-

ifelse(is.na(observables$NumberAttempts[is.na(observables$TrophyLevel)]),NA,"quit")
observables$TrophyLevel <- ordered(observables$TrophyLevel,
                                c("quit","silver","gold"))
```

### Multiple Attempts
Problem. What to do with multiple attempts. Two choices, take First, and Take

```

Last

```
```{r pivotObs}
obsFirst <-
  pivot_wider(observables, id_cols=uid,
              names_from = context,
              values_from = all_of(obsnames),
              values_fn = function(x) first(x,na_rm=TRUE))
obsLast <-
  pivot_wider(observables, id_cols=uid,
              names_from = context,
              values_from = all_of(obsnames),
              values_fn = function(x) last(x,na_rm=TRUE))
obsMax <-
  pivot_wider(observables, id_cols=uid,
              names_from=context,
              values_from=TrophyLevel,
              values_fn = function(x) max(x,na.rm=FALSE))
```
```

A lot of N/A observables, so we have total NA columns. Drop these.

```
```{r dropAllNA}
obsFirst<- obsFirst[,sapply(obsFirst,function(c) !all(is.na(c)))]
obsFirst <-
  obsFirst[,sapply(obsFirst,function(c) !is.numeric(c) || sum(c,na.rm=TRUE) != 0)]
obsLast<- obsLast[,sapply(obsLast,function(c) !all(is.na(c)))]
obsLast<-
  obsLast[,sapply(obsLast,function(c) !is.numeric(c) || sum(c,na.rm=TRUE) != 0)]
obsLast
```
### Level names
```

We currently don't need this, but might later.

```
```{r levelNames}
levels <-
gsub("TrophyLevel_(.*)", "\\1",grep("TrophyLevel_",names(obsFirst),value=TRUE))
levels
```
```

### Add Trophy Counts

```
```{r TrophyCounting}
obsMax %>% select(!uid) %>% as.matrix() %>% apply(1, function(row)
  table(ordered(row, c("quit","silver","gold")))) %>% t() -> tMax
obsMax <- cbind(obsMax,tMax)
obsFirst %>% select(ends_with("Trophies")) %>% as.matrix() %>% apply(1,
function(row)
  table(ordered(row, c("quit","silver","gold")))) %>% t() -> tFirst
obsFirst <- cbind(obsFirst,tFirst)
obsLast %>% select(ends_with("Trophies")) %>% as.matrix() %>% apply(1,
function(row)
  table(ordered(row, c("quit","silver","gold")))) %>% t() -> tLast
obsLast <- cbind(obsLast,tLast)
```
```

```

#### Write it out
```{r writeObs}
write_sheet(obsFirst,outID,sheet="ObsFirst")
write_sheet(obsLast,outID,sheet="ObsLast")
write_sheet(obsMax,outID,sheet="ObsMax")
write_csv(obsFirst,"data/PPIESFall2022obsFirst.csv")
write_csv(obsLast,"data/PPIESFall2022obsLast.csv")
write_csv(obsMax,"data/PPIESFall2022obsMax.csv")
```

## Learning Supports

Key is (uid,context,onWhat). Value is LS_duration. Can summarize with sum.

```{r cleanLS}
learningSupports <-
read_csv("https://pluto.coe.fsu.edu/Proc4/dongle/data/ppLS.BigStudy.csv")
learningSupports <-
  mutate(learningSupports,duration=LS_duration) %>%
  select(!any_of(c("...1","appid","LS_duration"))) %>%
  filter(!is.na(onWhat))
```

```{r lsPivot}
lsWide <-
  pivot_wider(learningSupports,
    id_cols=uid,
    names_from=c(context,onWhat),
    values_from=c(timestamp,duration),
    names_glue="{context}_{onWhat}_{.value}",
    values_fn=list(timestamp=~min(.x, na.rm=TRUE),
      duration=~ sum(.x,na.rm=TRUE)))
lsWide
```

Kill combinations that never occur.
```{r rmNAs}
all(sapply(lsWide,function(c) !all(is.na(c))))
```

```{r writeLS}
write_sheet(lsWide,outID,"LearningSupport")
write_csv(lsWide,"data/PPIESFall2022ls.csv")
```

## Join and Write

```{r BigJoin}
FSUSFall2022BigDaddy <- select(PPIDs,!Number) %>%
  full_join(pretest,by="StudyID",suffix=c("", ".pre")) %>%
  full_join(ecttest,by="StudyID",suffix=c("", ".ect")) %>%
  full_join(pottest,by="StudyID",suffix=c("", ".pot")) %>%
  left_join(BNscores,by=join_by(StudyID==uid),suffix=c("", ".bn")) %>%
  left_join(obsFirst,by=join_by(StudyID==uid),suffix=c("", ".first")) %>%
  left_join(obsLast,by=join_by(StudyID==uid),suffix=c("", ".last")) %>%
  left_join(obsMax,by=join_by(StudyID==uid),suffix=c("", ".max")) %>%

```

```
left_join(lsWide,by=join_by(StudyID==uid),suffix=c("", ".ls"))
```

```

```
{r writeBigDaddy}
write_sheet(FSUSFall2022BigDaddy,outID,"BigDaddy")
write_csv(FSUSFall2022BigDaddy,"data/PPIESFall2022Full.csv")
```

```

## SPSS Outpt

```
{r savOutput}
names(FSUSFall2022BigDaddy) <- gsub("-", "_",names(FSUSFall2022BigDaddy)) %>%
  gsub(" (in seconds)","",.,fixed=TRUE) %>%
  gsub(" ","",.) %>% gsub("(enter)","",.,fixed=TRUE) %>%
  gsub("Yippie!","Yippie",.,fixed=TRUE)
haven::write_sav(FSUSFall2022BigDaddy,"FSUSFall2022BigDaddy.sav")
```

```

## Metadata Export

```
{r pretestMetadata}
prechoices <- sapply(preSQ, function(sq) {
  if (length(sq$Choices) >0L) {
    choices <- sapply(sq$Choices,function(ch) ch$Display)
    c(choices,rep(NA_character_,8-length(choices)))
  } else {
    rep(NA_character_,8)
  }
})
prechoices <- t(prechoices)
colnames(prechoices) <- paste(1:ncol(prechoices))
```

```

```
{r pretestMeta2}
premeta <- data.frame(
  name=names(preSQ),
  qid=sapply(preSQ,function(sq) sq$QuestionID),
  qtag=sapply(preSQ,function(sq) sq$DataExportTag),
  text=sapply(preSQ,function(sq) sq$QuestionText),
  description=sapply(preSQ,function(sq) {
    if (is.null(sq$QuestionDescription))
      ""
    else
      sq$QuestionDescription
  }),
  choices=prechoices
)
premeta
```

```

### ECT

IMI meta-data is completely different from the rest, so needs special handling.

```
{r ecttestMetadata}
ectchoices <- sapply(ectSQ[names(ectSQ)!="IMI"], function(sq) {
  if (length(sq$Choices) >0L) {
```

```

    choices <- sapply(sq$Choices,function(ch) ch$Display)
    c(choices,rep(NA_character_,8-length(choices)))
  } else {
    rep(NA_character_,8)
  }
})
ectchoices <- t(ectchoices)
colnames(ectchoices) <- paste(1:ncol(ectchoices))

...

```{r ecttestMeta2}
ectmeta <- data.frame(
  name=paste(names(ectSQ[names(ectSQ)!="IMI"]), "ect", sep="."),
  qid=sapply(ectSQ[names(ectSQ)!="IMI"],function(sq) sq$QuestionID),
  qtag=sapply(ectSQ[names(ectSQ)!="IMI"],function(sq) sq$DataExportTag),
  text=sapply(ectSQ[names(ectSQ)!="IMI"],function(sq) sq$QuestionText),
  description=sapply(ectSQ[names(ectSQ)!="IMI"],function(sq) {
    if (is.null(sq$QuestionDescription))
      ""
    else
      sq$QuestionDescription
  }),
  choices=ectchoices
)

ectmeta
...

```{r IMIMeta}
IMISQ <- ectSQ[["IMI"]]
IMI.options <- c(sapply(IMISQ$Answers,function(ch) ch$Display),"8"=NA_character_)

imimeta <- data.frame(
  name=names(IMI.stems),
  qid=IMISQ$QuestionID,
  qtag=IMISQ$DataExportTag,
  text=IMI.stems,
  description=IMI.stems,
  choices=t(replicate(length(IMI.stems),IMI.options))
)
imimeta
...

### POT metadata

```{r pottestMetadata}
potchoices <- sapply(potSQ[names(potSQ)!="IMI"], function(sq) {
  if (length(sq$Choices) > 0L) {
    choices <- sapply(sq$Choices,function(ch) ch$Display)
    c(choices,rep(NA_character_,8-length(choices)))
  } else {
    rep(NA_character_,8)
  }
}

```

```

}))
potchoices <- t(potchoices)
colnames(ectchoices) <- paste(1:ncol(ectchoices))

```

```

```

```

```

```{r pottestMeta2}
potmeta <- data.frame(
  name=paste(names(potSQ[names(potSQ)!="IMI"]), "pot", sep="."),,
  qid=sapply(potSQ[names(potSQ)!="IMI"],function(sq) sq$QuestionID),
  qtag=sapply(potSQ[names(potSQ)!="IMI"],function(sq) sq$DataExportTag),
  text=sapply(potSQ[names(potSQ)!="IMI"],function(sq) sq$QuestionText),
  description=sapply(potSQ[names(potSQ)!="IMI"],function(sq) {
    if (is.null(sq$QuestionDescription))
      ""
    else
      sq$QuestionDescription
  }),
  choices=potchoices
)

```

```

potmeta

```

```

```

```

```

### Form B metadata

```

```

```{r postBMetadata}
bchoices <- sapply(postBSQ[names(postBSQ)!="IMI"], function(sq) {
  if (length(sq$Choices) > 0L) {
    choices <- sapply(sq$Choices,function(ch) ch$Display)
    c(choices,rep(NA_character_,8-length(choices)))
  } else {
    rep(NA_character_,8)
  }
})
bchoices <- t(bchoices)
colnames(bchoices) <- paste(1:ncol(bchoices))

```

```

```

```

```

```{r postBMeta2}
bmeta <- data.frame(
  name=paste(names(postBSQ[names(postBSQ)!="IMI"]), "pot", sep="."),,
  qid=sapply(postBSQ[names(postBSQ)!="IMI"],function(sq) sq$QuestionID),
  qtag=sapply(postBSQ[names(postBSQ)!="IMI"],function(sq) sq$DataExportTag),
  text=sapply(postBSQ[names(postBSQ)!="IMI"],function(sq) sq$QuestionText),
  description=sapply(postBSQ[names(postBSQ)!="IMI"],function(sq) {
    if (is.null(sq$QuestionDescription))
      ""
    else
      sq$QuestionDescription
  }),
  choices=bchoices
)

```

```
bmeta
```
```

```
### Write it out
```

```
` `{r writemeta}
write_sheet(rbind(premeta,ectmeta,imimeta,potmeta),outID,"Metadata")
` `
```