wx3 analysis

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```
rm(list=ls())
library(ggplot2)
library(reshape2)
library(plyr)
library(bootstrap)
setwd("~/github/local/pragmods")
#source("useful.R")
d <- read.csv("seq_data/pragmods_wx3.anondata.csv")</pre>
\#head(d)
```

```
Exclude participants that either failed manipulation check or were rejected.
exclude <- d$assignmentstatus == "Rejected" |
  d$Answer.name_check_correct == "\"FALSE\""
sum(exclude)
## [1] 3
mean(exclude)
## [1] 0.06
d <- subset(d, exclude == FALSE)</pre>
d$Answer.choice_correct_1 <- factor(as.logical(d$Answer.choice_correct_1))
d$Answer.choice_correct_2 <- factor(as.logical(d$Answer.choice_correct_2))
d$Answer.choice_correct_3 <- factor(as.logical(d$Answer.choice_correct_3))
d$Answer.choice_correct_4 <- factor(as.logical(d$Answer.choice_correct_4))
d$Answer.choice_correct_5 <- factor(as.logical(d$Answer.choice_correct_5))
d$Answer.choice_correct_6 <- factor(as.logical(d$Answer.choice_correct_6))
```

Restructure data to have trial variable. (working around mturk data submission limitations; can't safely submit arrays, so we have to use separate variables for each trial)

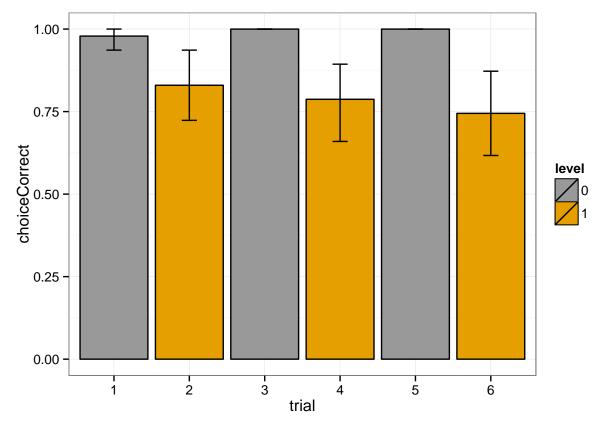
```
d$level_1 <- factor(rep(0,nrow(d)))
d$level_2 <- factor(rep(1,nrow(d)))
d$level_3 <- factor(rep(0,nrow(d)))</pre>
d$level_4 <- factor(rep(1,nrow(d)))
d$level_5 <- factor(rep(0,nrow(d)))</pre>
d$level_6 <- factor(rep(1,nrow(d)))
trial.df <- function(tn){</pre>
  df <- data.frame(id = d$workerid,</pre>
                    seqCond = d$Answer.sequence condition,
                    trial = factor(rep(tn,nrow(d))),
```

```
level = d[sprintf("level_%d",tn)],
                   item = d[sprintf("Answer.item_%d",tn)],
                   targetProp = d[sprintf("Answer.target_prop_%d",tn)],
                   distractorProp = d[sprintf("Answer.distractor_prop_%d",tn)],
                   targetPosition = d[sprintf("Answer.target_position_%d",tn)],
                   distractorPosition = d[sprintf("Answer.distractor_position_%d",tn)],
                   choice = d[sprintf("Answer.choice_%d",tn)],
                   choiceCorrect = d[sprintf("Answer.choice correct %d",tn)])
  names(df) <- c("id", "seqCond", "trial", "level", "item", "targetProp", "distractorProp",</pre>
                 "targetPosition", "distractorPosition", "choice", "choiceCorrect")
  return(df)
}
d2 <- rbind(trial.df(1),trial.df(2),trial.df(3),trial.df(4),trial.df(5),trial.df(6))
summary(d2)
                         {\tt seqCond}
##
          id
                                     trial level
                                                                   item
## Min.
          : 1.0
                   "(0w1w)x3":282
                                                    "boat"
                                     1:47
                                            0:141
                                                                     :66
## 1st Qu.:12.0
                                     2:47
                                            1:141
                                                    "Christmas tree":54
## Median :24.0
                                     3:47
                                                    "friend"
                                                                     :30
## Mean :25.2
                                     4:47
                                                    "pizza"
                                                                     :54
## 3rd Qu.:39.0
                                                    "snowman"
                                     5:47
                                                                     :48
## Max.
         :50.0
                                     6:47
                                                    "sundae"
                                                                     :30
##
##
                          distractorProp targetPosition distractorPosition
          targetProp
                                 : 24
                                          "left" : 78
## "motor"
              : 24
                      "motor"
                                                          "left" : 78
                      "mushrooms": 24
## "mushrooms": 24
                                          "middle":108
                                                          "middle":108
## "sail"
              : 24
                      "sail"
                                 : 24
                                          "right" : 96
                                                          "right": 96
## "hat"
               : 21
                      "hat"
                                  : 21
## "cabin"
               : 18
                      "cabin"
                                  : 18
## "lights"
                      "lights"
               : 18
                                  : 18
## (Other)
               :153
                      (Other)
                                  :153
##
                    choiceCorrect
          choice
## "logical": 25
                    FALSE: 31
## "target" :251
                    TRUE: 251
## "foil"
           : 6
##
##
##
##
#statistics for boolean factors; copied from useful.R, with a slightly different mean function to work
1.mean <- function(...){mean(as.logical(...))}</pre>
1.theta <- function(x,xdata,na.rm=T) {1.mean(xdata[x],na.rm=na.rm)}</pre>
1.ci.low <- function(x,na.rm=T) {</pre>
  1.mean(x,na.rm=na.rm) - quantile(bootstrap(1:length(x),1000,1.theta,x,na.rm=na.rm)$thetastar,.025,na.
1.ci.high <- function(x,na.rm=T) {</pre>
  quantile(bootstrap(1:length(x),1000,1.theta,x,na.rm=na.rm)$thetastar,.975,na.rm=na.rm) - 1.mean(x,na.rm=na.rm)
ms <- aggregate(choiceCorrect ~ trial + level,data = d2,1.mean)
ms$cil <- aggregate(choiceCorrect ~ trial + level, data = d2, 1.ci.low)$choiceCorrect
ms$cih <- aggregate(choiceCorrect ~ trial + level, data = d2, l.ci.high)$choiceCorrect
```

```
#colorblind-friendly color palettes
cbPalette <- c("#999999", "#E69F00", "#56B4E9", "#009E73", "#F0E442", "#0072B2", "#D55E00", "#CC79A7")
cbbPalette <- c("#000000", "#E69F00", "#56B4E9", "#009E73", "#F0E442", "#0072B2", "#D55E00", "#CC79A7")</pre>
```

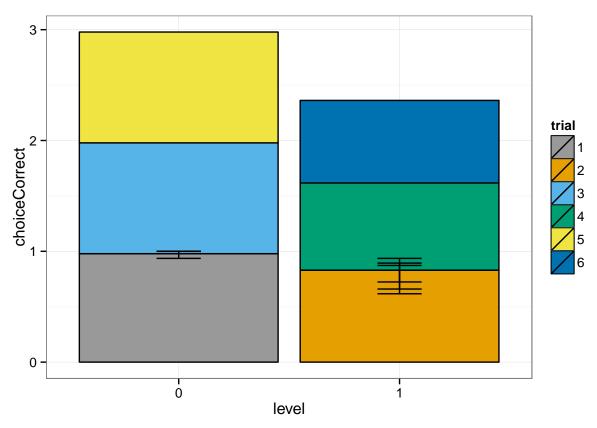
Compare performance on trials, in chronological order.

```
ggplot(data = ms, aes(x = trial, y = choiceCorrect, fill = level)) +
  geom_bar(stat = "identity", color = "black") +
  geom_errorbar(aes(ymin = choiceCorrect - cil, ymax = choiceCorrect + cih), width = 0.2) +
  #facet_grid(. ~ seqCond) +
  theme_bw() +
  scale_fill_manual(values=cbPalette)
```



Compare performance on level-0 vs level-1 trials.

```
ggplot(data = ms, aes(x = level, y = choiceCorrect, fill = trial)) +
geom_bar(stat = "identity", color = "black") +
geom_errorbar(aes(ymin = choiceCorrect - cil, ymax = choiceCorrect + cih), width = 0.2) +
#facet_grid(. ~ seqCond) +
theme_bw() +
scale_fill_manual(values=cbPalette)
```



Chi-squared test between level-1 inferences (TODO...)

```
## seqCond foil target logical
## 1 "(0w1w)x3" 6 111 24
```

#chisq.test(cont_table[1:2,2:4])