seq_analysis

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July 28, 2014

```
rm(list=ls())
library(ggplot2)
library(reshape2)
library(plyr)
library(bootstrap)
#source("useful.R")
d <- read.csv("pragmods_seq.results.tsv",sep="\t")
#head(d)</pre>
```

Exclude participants that either failed manipulation check or were rejected.

```
exclude <- d$assignmentstatus == "Rejected" |
  d$Answer.name_check_correct == "\"FALSE\""
sum(exclude)</pre>
```

[1] 4

```
mean(exclude)
```

[1] 0.04

```
d <- subset(d, exclude == FALSE)

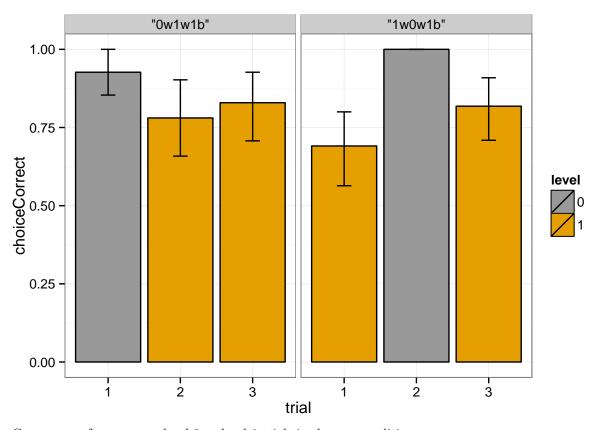
d$Answer.choice_correct_1 <- factor(as.logical(d$Answer.choice_correct_1))
d$Answer.choice_correct_2 <- factor(as.logical(d$Answer.choice_correct_2))
d$Answer.choice_correct_3 <- factor(as.logical(d$Answer.choice_correct_3))</pre>
```

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)

```
#statistics for boolean factors; copied from useful.R, with a slightly different mean function to work
l.mean <- function(...){mean(as.logical(...))}
l.theta <- function(x,xdata,na.rm=T) {l.mean(xdata[x],na.rm=na.rm)}
l.ci.low <- function(x,na.rm=T) {
   l.mean(x,na.rm=na.rm) - quantile(bootstrap(1:length(x),1000,l.theta,x,na.rm=na.rm)$thetastar,.025,na.d.
l.ci.high <- function(x,na.rm=T) {
   quantile(bootstrap(1:length(x),1000,l.theta,x,na.rm=na.rm)$thetastar,.975,na.rm=na.rm) - l.mean(x,na.d.
ms <- aggregate(choiceCorrect ~ seqCond + trial + level,data = d2,l.mean)
ms$cil <- aggregate(choiceCorrect ~ seqCond + trial + level, data = d2, l.ci.low)$choiceCorrect
ms$cih <- aggregate(choiceCorrect ~ seqCond + trial + level, data = d2, l.ci.high)$choiceCorrect
#colorblind-friendly color palettes
cbPalette <- c("#999999", "#E69F00", "#56B4E9", "#009E73", "#F0E442", "#0072B2", "#D55E00", "#CC79A7")
cbbPalette <- c("#0000000", "#E69F00", "#56B4E9", "#009E73", "#F0E442", "#0072B2", "#D55E00", "#CC79A7")</pre>
```

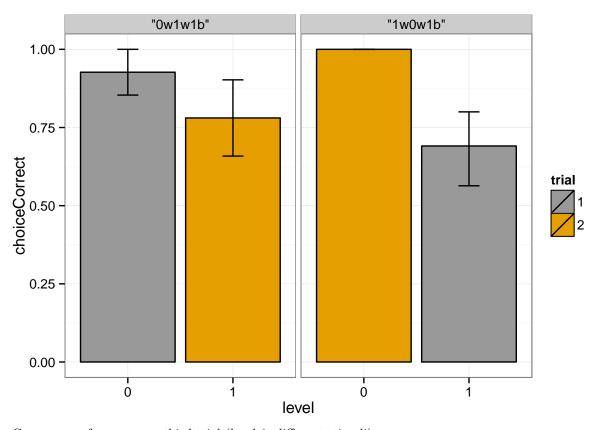
Compare performance on trials, in chronological order; facet by ordering condition.

```
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 in the two conditions

```
ggplot(data = subset(ms,trial != "3"), 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)
```



Compare performance on third trial (level-1, different stimuli)

```
ggplot(data = subset(ms,trial=="3"), aes(x = seqCond, y = choiceCorrect, fill = seqCond)) +
  geom_bar(stat = "identity", color = "black") +
  geom_errorbar(aes(ymin = choiceCorrect - cil, ymax = choiceCorrect + cih), width = 0.2) +
  theme_bw() +
  scale_fill_manual(values=cbPalette)
```

