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# R course for beginners
# Week 7
# assignment by Noam Gabay, id 208540815
### preprocessing ----
# creating raw data
subjects = dir("stroop data")
df = data.frame()
for (subject in subjects)
 temp data = read.csv(file.path("stroop data", subject))
  df = rbind(df, temp_data)
# piping
library(dplyr)
df = df \mid >
 mutate(
    task = ifelse(grepl("word reading", condition), "word reading",
                  ifelse(grepl("ink_naming", condition), "ink_naming", NA)),
    congruency = ifelse(grepl(" cong", condition), "congruent",
                        ifelse(grepl("incong", condition), "incongruent", NA)),
  )
df = df \mid >
 mutate(
   accuracy = (participant response == correct response) * 1
df = df \mid >
  select(subject, task, congruency, block, trial, accuracy, rt)
df = df \mid >
 mutate(
    subject = factor(subject),
    task = factor(task, levels = c("word reading", "ink naming")),
    congruency = factor(congruency, levels = c("congruent", "incongruent")),
save(df, file = "raw data.rdata")
### creating filtered data ----
# number of subjects
n subjects = length(unique(df$subject))
# drop rows with NAs
df filtered = na.omit(df)
# condition for rt (assuming rt is in ms)
df filtered = df filtered |>
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filter(
    (df filtered$rt > 300) & (df filtered$rt < 3000)
# trial percentage
n trials original = df |>
  group by (subject) |>
  summarize(trials = n())
n trials left = df filtered |>
  group by(subject) |>
  summarize(trials = n())
trial data = data.frame(
  subject = unique(df filtered$subject),
  percent remaining = n trials left$trials / n trials original$trials * 100 ,
  percent_removed = (n_trials_original$trials - n_trials_left$trials) /
n trials original$trials *100
  )
print(trial data)
trial summary = trial data |>
  summarize(
   mean removed = mean(percent removed, na.rm = TRUE),
    sd removed = sd(percent removed, na.rm = TRUE)
print(trial_summary)
save(df filtered, file= "filtered data.csv")
### descriptive statistics ----
summary conditions = df filtered |>
  group by (task, congruency) |>
  summarize(
    mean rt = mean(rt, na.rm = TRUE),
    sd rt = sd(rt, na.rm = TRUE),
   mean accuracy = mean(accuracy, na.rm = TRUE),
    sd accuracy = sd(accuracy, na.rm = TRUE)
### plotting ----
library(ggplot2)
#plot rt
ggplot(summary\_conditions, aes(x = task, y = mean\_rt, fill = congruency)) +
  geom bar(stat = "identity", position = position dodge()) +
  geom errorbar(aes(ymin = mean rt - sd rt, ymax = mean rt + sd rt),
                position = position dodge(0.9), width = 0.25) +
    title = "Mean Reaction Times by Task and Congruency",
    x = "Task",
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y = "Mean Reaction Time (ms)",
   fill = "Congruency"
  ) +
  theme minimal()
# scatter plot
ggplot(df filtered, aes(x = task, y = rt, color = congruency)) +
  geom point(alpha = 0.5, position = position jitter(width = 0.2)) + # נקודות עם
רנדומליות קלה
  labs(
    title = "Reaction Times by Task and Congruency",
    x = "Task",
    y = "Reaction Time (ms)",
   color = "Congruency"
  ) +
  theme_minimal()
### stats ----
library(lme4)
# to account for the fact that each subject did all conditions
# I consulted with chatGPT on how to include subject as a random effect
# and got:
# (1 | subject) for variance in intercept between participants
# (task * congruency | subject) for variance in slope between participants
model = lmer(rt \sim task * congruency + (1 + task * congruency | subject), data =
df filtered)
coef(model)
summary(model)
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