Lab 9

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Problem 1

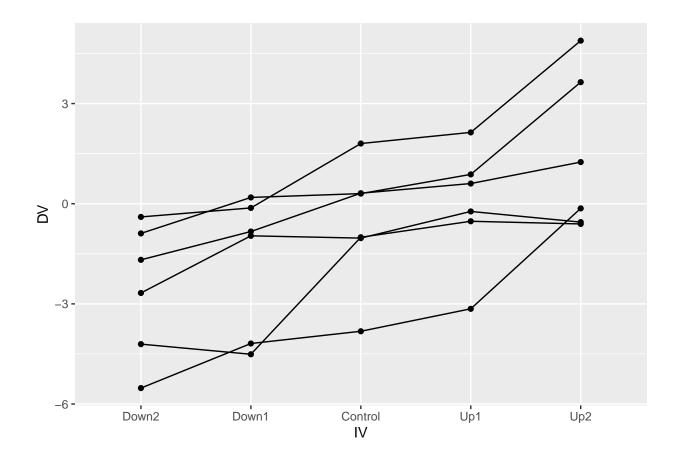
Create an R script that can generate simulated data for the following repeated measures design. (2 points) A. The dependent variable is assumed to come from a normal distribution with mean = 0 and standard deviation = 1.

- B. There is one repeated measures factor with 5 levels (Down1, Down2, Control, Up1, Up2). The control group is assumed to have no effect. The Down1 and Down2 levels shift the mean down by 1 and 2 standard deviations, respectively. The Up1 and Up2 levels shift the mean up by 1 and 2 standard deviations, respectively.
- C. There are 6 subjects in the experiment, and they are each measured once in each condition. The 6 subjects are assumed to be different from one another (e.g., they will have different baseline means in the control condition), but they will all be influenced by the IV in the exact same way (e.g., no interaction).

```
sim_data <- tibble(
subjects = rep(1:6, each = 5),
IV = rep(c("Down2", "Down1", "Control", "Up1", "Up2"),6),
DV = rnorm(6*5,c(-2,-1,0,1,2),1)
)%>% mutate(DV = DV + rep(2:-3, each = 5))

#change order of x-axis, save this code
sim_data$IV <- factor(sim_data$IV, levels = c("Down2", "Down1", "Control", "Up1", "Up2"))
sim_data$subjects <- as.factor(sim_data$subjects)

ggplot(sim_data, aes(x=IV, y= DV, group = subjects))+
geom_point()+
geom_line()</pre>
```



Problem 2

Run a simulation to determine the proportion of experiments that would return a significant result for the above design. Assume that the effect of the levels of the IV are increments of .1 of a standard deviation, rather than increments of 1 as in the above design.

```
save_p <- c()
for (i in 1:1000){

sim_data <- tibble(
subjects = rep(1:6, each = 5),
    IV = rep(c("Down2", "Down1", "Control", "Up1", "Up2"),6),
    DV = rnorm(6*5,c(-.2,-.1,0,.1,.2),1)
) %>% mutate(DV = DV + rep(rnorm(6, 0, 1), each = 5))

sim_data$IV <- factor(sim_data$IV, levels = c("Down2", "Down1", "Control", "Up1", "Up2"))

sim_data$subjects <- as.factor(sim_data$subjects)

aov_out <- aov(DV ~ IV + Error(subjects), sim_data) %>% summary
save_p[i] <- aov_out[2]$`Error: Within`[[1]]$`Pr(>F)`[1]
}

length(save_p[save_p < .05])/1000</pre>
```

Problem 3

Demonstrate that the Godden and Baddeley example data from the textbook (19.5), which used a 2x2 repeated measures design, can be be analyzed with one-sample t-tests to return the same results. Specifically, show the one-sample t-tests for each main effect and the interaction. (2 points)

```
godden baddeley <- tribble(~Subjects,~LearningPlace,~TestingPlace,~Recall,
    "s1","On Land","On Land",34,
    "s2","On Land","On Land",37,
    "s3","On Land","On Land",27,
    "s4","On Land","On Land",43,
    "s5","On Land","On Land",44,
    "s1","On Land","Under Sea",18,
    "s2","On Land","Under Sea",21,
    "s3","On Land","Under Sea",25,
    "s4","On Land","Under Sea",37,
    "s5","On Land","Under Sea",34,
    "s1","Under Sea","On Land",14,
    "s2","Under Sea","On Land",21,
    "s3","Under Sea","On Land",31,
    "s4","Under Sea","On Land",27,
    "s5","Under Sea","On Land",32,
    "s1","Under Sea","Under Sea",22,
    "s2","Under Sea","Under Sea",25,
    "s3","Under Sea","Under Sea",33,
    "s4","Under Sea","Under Sea",33,
    "s5","Under Sea","Under Sea",42
    )
godden baddeley <- godden baddeley %>%
 mutate(Subjects = as.factor(Subjects),
     LearningPlace = as.factor(LearningPlace),
     TestingPlace = as.factor(TestingPlace))
```

Main Effect: Learning Place

```
lp_mean <- godden_baddeley %>%
  group_by(Subjects, LearningPlace) %>%
  summarize(mean_recall = mean(Recall))

## `summarise()` regrouping output by 'Subjects' (override with `.groups` argument)

t.test(mean_recall ~ LearningPlace, paired = TRUE, data = lp_mean)

## Paired t-test
##
```

```
## data: mean_recall by LearningPlace
## t = 1.4142, df = 4, p-value = 0.2302
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -3.852973 11.852973
## sample estimates:
## mean of the differences
## 4
```

Main Effect: Testing Place

```
tp mean <- godden baddeley %>%
group by (Subjects, TestingPlace) %>%
summarize(mean recall = mean(Recall))
## 'summarise()' regrouping output by 'Subjects' (override with '.groups' argument)
t.test(mean_recall ~ TestingPlace, paired = TRUE, data = tp_mean)
##
## Paired t-test
## data: mean recall by TestingPlace
## t = 1.5811, df = 4, p-value = 0.189
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -1.511956 5.511956
## sample estimates:
## mean of the differences
##
                2
```

Interaction

```
filter(LearningPlace == "Under Sea",
    TestingPlace == "Under Sea") %>%

pull(Recall)

t.test((LL - LS) - (SL - SS), mu = 0)

##

## One Sample t-test
##

## data: (LL - LS) - (SL - SS)
## t = 4.4721, df = 4, p-value = 0.01106
## alternative hypothesis: true mean is not equal to 0

## 95 percent confidence interval:
## 6.066688 25.933312
## sample estimates:
```

mean of x

16

##