

HW-Week 7

Md Ariful Haque Miah

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Problem 4.3

```
library(GAD)
```

```
## Loading required package: matrixStats
```

```
## Loading required package: R.methodsS3
```

```
## R.methodsS3 v1.8.2 (2022-06-13 22:00:14 UTC) successfully loaded. See ?R.methodsS3 for help.
```

```
chemical<-c(rep(1,5),rep(2,5),rep(3,5),rep(4,5))
```

```
bolt<-c(rep(seq(1,5),4))
```

```
obs <- c(73,68,74,71,67,73,67,75,72,70,75,68,78,73,68,73,71,75,75,69)
```

```
chemical<-as.fixed(chemical)
```

```
bolt<-as.fixed(bolt)
```

```
obs <- as.numeric(obs)
```

```
dat <- data.frame(chemical,bolt,obs)
```

```
dat
```

```
##      chemical bolt obs
## 1          1     1  73
## 2          1     2  68
## 3          1     3  74
## 4          1     4  71
## 5          1     5  67
## 6          2     1  73
## 7          2     2  67
## 8          2     3  75
## 9          2     4  72
## 10         2     5  70
## 11         3     1  75
## 12         3     2  68
## 13         3     3  78
## 14         3     4  73
## 15         3     5  68
## 16         4     1  73
## 17         4     2  71
## 18         4     3  75
## 19         4     4  75
## 20         4     5  69
```

```
str(dat)
```

```
## 'data.frame':   20 obs. of  3 variables:
```

```
## $ chemical: Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 2 2 2 2 2 ...
```

```
## $ bolt    : Factor w/ 5 levels "1","2","3","4",...: 1 2 3 4 5 1 2 3 4 5 ...
```

```
## $ obs      : num  73 68 74 71 67 73 67 75 72 70 ...
```

```
model<-lm(obs~chemical+bolt,data=dat)
gad(model)
```

```
## Analysis of Variance Table
```

```
##
```

```
## Response: obs
```

```
##           Df Sum Sq Mean Sq F value    Pr(>F)
## chemical  3  12.95   4.317   2.3761    0.1211
## bolt      4 157.00  39.250 21.6055 2.059e-05 ***
## Residual 12   21.80   1.817
```

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Hypothesis:

$H_0 : \tau_i = 0$ for all i

$H_a : \tau_i \neq 0$ for some i

Model Equation:

$$y_{ij} = \mu + \tau_i + \beta_j + e_{ij}$$

where μ = grand mean, τ_i = Chemical effect, β_j = Bolt (block), e_{ij} = random error $\sim N(0, \sigma^2)$

P value (0.1211) $> \alpha$ (0.05) hence we fail to reject H_0 .

There is no difference among the four chemical agents at $\alpha = 0.05$ level.

Problem 4.16

Calculate model parameters τ_i : Formula: $\tau_i = \mu_i - \mu$

```
obs <- c(73,68,74,71,67,73,67,75,72,70,75,68,78,73,68,73,71,75,75,69)
mean(obs)
```

```
## [1] 71.75
```

```
tau_1 <- mean(obs[1:5])-mean(obs)
tau_1
```

```
## [1] -1.15
```

```
tau_2 <- mean(obs[6:10])-mean(obs)
tau_2
```

```
## [1] -0.35
```

```
tau_3 <- mean(obs[11:15])-mean(obs)
tau_3
```

```
## [1] 0.65
```

```
tau_4 <- mean(obs[16:20])-mean(obs)
tau_4
```

```
## [1] 0.85
```

$\tau_1 = -1.15$, $\tau_2 = -0.35$, $\tau_3 = 0.65$, $\tau_4 = 0.85$

Calculate model parameters β_j : Formula: $\beta_j = \mu_j - \mu$

```
obs <- c(73,68,74,71,67,73,67,75,72,70,75,68,78,73,68,73,71,75,75,69)
beta_1 <- mean(c(73,73,75,73))-mean(obs)
beta_1
```

```
## [1] 1.75
```

```
beta_2 <- mean(c(68,67,68,71))-mean(obs)
beta_2
```

```
## [1] -3.25
```

```
beta_3 <- mean(c(74,75,78,75))-mean(obs)
beta_3
```

```
## [1] 3.75
```

```
beta_4 <- mean(c(71,72,73,75))-mean(obs)
beta_4
```

```
## [1] 1
```

```
beta_5 <- mean(c(67,70,68,69))-mean(obs)
beta_5
```

```
## [1] -3.25
```

$\beta_1 = 1.75, \beta_2 = -3.25, \beta_3 = 3.75, \beta_4 = 1, \beta_5 = -3.25$

Problem 4.22

```
library(GAD)
batch <- c(rep(1,5),rep(2,5),rep(3,5),rep(4,5),rep(5,5))
day <- c(rep(seq(1,5),5))
ingr <- c("A","B","D","C","E",
          "C","E","A","D","B",
          "B","A","C","E","D",
          "D","C","E","B","A",
          "E","D","B","A","C")
obs <- c(8,7,1,7,3,
        11,2,7,3,8,
        4,9,10,1,5,
        6,8,6,6,10,
        4,2,3,8,8)
batch <- as.fixed(batch)
day <- as.fixed(day)
ingr <- as.fixed(ingr)
obs <- as.numeric(obs)
data <- data.frame(batch,day,ingr,obs)
data
```

```
##      batch day ingr obs
## 1      1    1   A    8
## 2      1    2   B    7
## 3      1    3   D    1
## 4      1    4   C    7
## 5      1    5   E    3
## 6      2    1   C   11
## 7      2    2   E    2
## 8      2    3   A    7
```

```
## 9      2    4    D    3
## 10     2    5    B    8
## 11     3    1    B    4
## 12     3    2    A    9
## 13     3    3    C   10
## 14     3    4    E    1
## 15     3    5    D    5
## 16     4    1    D    6
## 17     4    2    C    8
## 18     4    3    E    6
## 19     4    4    B    6
## 20     4    5    A   10
## 21     5    1    E    4
## 22     5    2    D    2
## 23     5    3    B    3
## 24     5    4    A    8
## 25     5    5    C    8
```

```
str(data)
```

```
## 'data.frame':   25 obs. of  4 variables:
## $ batch: Factor w/ 5 levels "1","2","3","4",...: 1 1 1 1 1 2 2 2 2 2 ...
## $ day : Factor w/ 5 levels "1","2","3","4",...: 1 2 3 4 5 1 2 3 4 5 ...
## $ ingr : Factor w/ 5 levels "A","B","C","D",...: 1 2 4 3 5 3 5 1 4 2 ...
## $ obs : num  8 7 1 7 3 11 2 7 3 8 ...
```

```
model <- aov(obs~batch+day+ingr, data=data)
summary(model)
```

```
##           Df Sum Sq Mean Sq F value    Pr(>F)
## batch      4  15.44     3.86    1.235 0.347618
## day        4  12.24     3.06    0.979 0.455014
## ingr       4 141.44    35.36   11.309 0.000488 ***
## Residuals 12   37.52     3.13
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Hypothesis H_0 : $\tau_i = 0$ for all i

H_a : $\tau_i \neq 0$ for some i , $i=1,2,3,4,5$

Model equation: $X_{ijk} = \mu + \tau_i + \beta_j + \alpha_k + e_{ijk}$

where, μ =grand mean, τ_i =Ingredients effect, β_j = Batch (block 1) α_k = Day (block 2), e_{ijk} = random error $\sim N(0, \sigma^2)$

Conclusions:

Hence the p value (0.000488) of ingredients is highly significant and less than $\alpha=0.05$ so we reject the null hypothesis H_0 .

Therefore the effect of five different ingredients (A,B,C,D,E) on the reaction time of a chemical process is significantly different.

Source Code

```
library(GAD)
chemical<-c(rep(1,5),rep(2,5),rep(3,5),rep(4,5))
bolt<-c(rep(seq(1,5),4))
obs <- c(73,68,74,71,67,73,67,75,72,70,75,68,78,73,68,73,71,75,75,69)
```

```

chemical<-as.fixed(chemical)
bolt<-as.fixed(bolt)
obs <- as.numeric(obs)
dat <- data.frame(chemical,bolt,obs)
dat
str(dat)
model<-lm(obs~chemical+bolt,data=dat)
gad(model)

obs <- c(73,68,74,71,67,73,67,75,72,70,75,68,78,73,68,73,71,75,75,69)
mean(obs)
tau_1 <- mean(obs[1:5])-mean(obs)
tau_1
tau_2 <- mean(obs[6:10])-mean(obs)
tau_2
tau_3 <- mean(obs[11:15])-mean(obs)
tau_3
tau_4 <- mean(obs[16:20])-mean(obs)
tau_4

obs <- c(73,68,74,71,67,73,67,75,72,70,75,68,78,73,68,73,71,75,75,69)
beta_1 <- mean(c(73,73,75,73))-mean(obs)
beta_1
beta_2 <- mean(c(68,67,68,71))-mean(obs)
beta_2
beta_3 <- mean(c(74,75,78,75))-mean(obs)
beta_3
beta_4 <- mean(c(71,72,73,75))-mean(obs)
beta_4
beta_5 <- mean(c(67,70,68,69))-mean(obs)
beta_5

library(GAD)
batch <- c(rep(1,5),rep(2,5),rep(3,5),rep(4,5),rep(5,5))
day <- c(rep(seq(1,5),5))
ingr <- c("A","B","D","C","E",
          "C","E","A","D","B",
          "B","A","C","E","D",
          "D","C","E","B","A",
          "E","D","B","A","C")
obs <- c(8,7,1,7,3,
        11,2,7,3,8,
        4,9,10,1,5,
        6,8,6,6,10,
        4,2,3,8,8)
batch <- as.fixed(batch)
day <- as.fixed(day)
ingr <- as.fixed(ingr)
obs <- as.numeric(obs)
data <- data.frame(batch,day,ingr,obs)
data
str(data)
model <- aov(obs~batch+day+ingr, data=data)

```

```
summary(model)
```