A4q3

Part a

The randomization performed by randomly assigning treatments to the bolt. Witch means we have ramdon chemical X and chemical y within each bolts.

Part b

$$Y_{ijk} = \mu + \alpha_i + \lambda_j + \tau_{ij} + \beta_k + R_{ijk}$$

where mu is the over all mean alpha is the hemical X effect lambda is the hemical Y effect tau is the interaction effect beta is the Bolt effect R is the random Error where i is 1 to ta, ta = 2(# of A); j is 1 to tb, tb = 2(# of B); k is 1 to r, r = 4(# of blcok) constraints:

$$\sum_{i=1}^{ta}\alpha_i=0, \sum_{j=1}^{tb}\lambda_j=0, \sum_{k=1}^{r}\beta_k=0, \sum_{i}\tau_{ij}=0, for \quad \forall i; \sum_{j}\tau_{ij}=0, for \quad \forall j$$

Part c

C) 1. Hypothesis Ho;
$$\theta = 0$$
Since $\theta = \tau_1 - \tau_2$

Test statistic

2. This; i diad all y_{11} , y_{12} , y_{21} , y_{32} by k code
$$\theta = -0.6 \text{ by } k \text{ code}$$

$$\theta^2 = 1.82 \text{ by } k \text{ code from } ml, \text{ since } \hat{\xi}^2 = \text{mean square of Residual mode.}$$

$$T - \text{obs} = \frac{1 - 0.6!}{\sqrt{1.82} \int_{\frac{1}{5}}^{2} + \frac{1}{5} + \frac{1}{5}!} \approx \frac{4.4470.4972}{\sqrt{1.82} \int_{\frac{1}{5}}^{2} + \frac{1}{5} + \frac{1}{5}!} \approx \frac{4.4470.4972}{\sqrt{1.82} \int_{\frac{1}{5}}^{2} + \frac{1}{5}!} \approx \frac{1.4470.4972}{\sqrt{1.82} \int_{\frac{1}{5}}^{2} + \frac{1}{5}!} \approx \frac$$

The code is showed in the next page.

```
> q3block =matrix(NA, nrow=4, ncol=5)
> q3block[1,] =c(73,68,74,71,67)
> q3block[2,] =c(73,67,75,72,70)
> q3block[3,] =c(75,68,78,73,68)
> q3block[4,] =c(73,71,75,75,69)
> rownames(q3block) =c('1','2','3','4')
> mu_11 = mean(q3block[1,])
> mu_12 = mean(q3block[2,])
> mu_21 = mean(q3block[3,])
> mu_22 = mean(q3block[4,])
> baths = c(q3block[1,],q3block[2,],q3block[3,],q3block[4,])
> treatment = rep(c(1:4), each = 5)
> treatment = factor(treatment)
> bolt = rep(c(1:5),4)
> bolt = factor(bolt)
> m1 = aov(baths~treatment+bolt)
> summary(m1)
           Df Sum Sq Mean Sq F value
                                     Pr(>F)
treatment
            3 12.95
                      4.32
                             2.376
                                     0.121
                      39.25 21.606 2.06e-05 ***
bolt
            4 157.00
Residuals 12 21.80
                       1.82
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> sigma_sq = 1.82
> theta_hat = (mu_11 - mu_12) - (mu_21 - mu_22)
> T_obs = abs(theta_hat) / (sqrt(sigma_sq) * sqrt(4/5))
> T_obs
[1] 0.4972452
> 2*pt(T_obs,df=12,lower.tail = FALSE)
[1] 0.6280016
  Part d
    (i) since the p-value 0.121 > 0.05 (from part(c) code), there is no evidence against HO
   (ii) since the p-value 0.6277 > 0.5, there is NO evidence against HO.
Code of part d:
> h_x = rep(c('low', 'high'), each = 10)
> h_y = rep(rep(c('low', 'high'), each = 5), 2)
> m2 = aov(baths \sim h_x * h_y + bolt)
> summary(m2)
               Df Sum Sq Mean Sq F value
                                                  Pr(>F)
                1 11.25
                              11.25
                                        6.193
                                                  0.0285 *
h_x
                     1.25
h_y
                               1.25
                                        0.688
                                                  0.4230
                1
bolt
                4 157.00
                              39.25 21.606 2.06e-05 ***
h_x:h_y
                1
                     0.45
                              0.45
                                        0.248
                                                 0.6277
Residuals
               12 21.80
                              1.82
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```