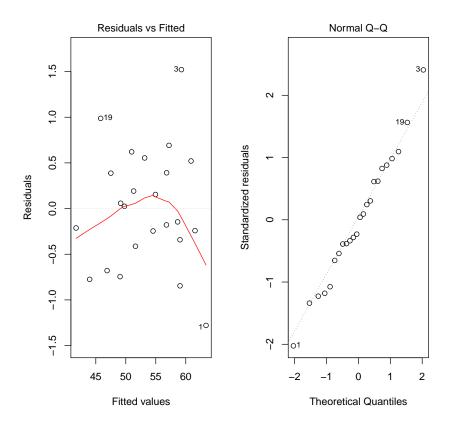
Stat 207 HW4

Cheng Luo 912466499 Fan Wu 912538518

February 2, 2015

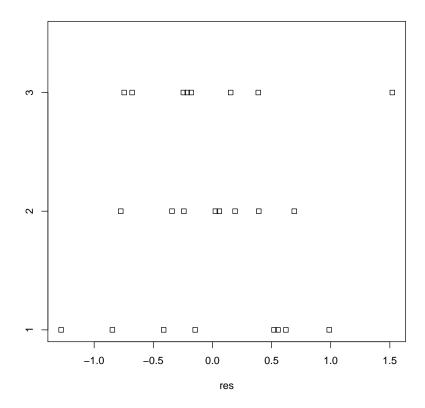


```
dat = read.table("CH27PR06.txt")
(a)
   names(dat) = c("Y", "S", "A")
   dat1 = dat
   dat$A = factor(dat$A)
   dat$S = factor(dat$S)
   r = length(unique(dat$S))
   a = length(unique(dat$A))
   model = aov(Y^{-}A + S, data = dat)
   res = resid(model)
   res
                    2
                             3
  ## -1.27916667 -0.24166667 1.52083333 -0.84583333 0.69166667 0.15416667
     7 8 9 10
                                        11
    ##
  ##
       13
              14 15
                               16
                                         17
                                                       18
  ## 0.52083333 -0.34166667 -0.17916667 -0.14583333 0.39166667 -0.24583333
     19 20 21
                               22
                                              23
  ## 0.98750000 -0.77500000 -0.21250000 -0.41250000 0.02500000 0.38750000
   par(mfrow = c(1,2))
   plot(model, which = 1)
  plot(model, which = 2)
```



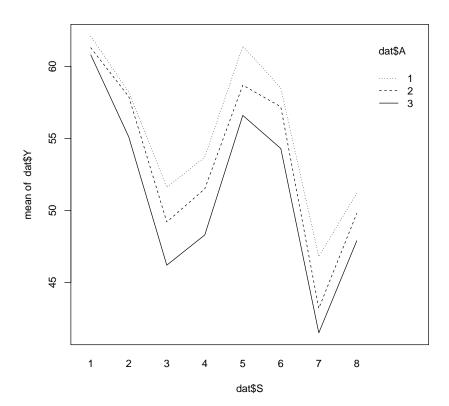
The residuals versus fitted values plots shows no sign for unequal variance. And the QQ-plot indicates approximately normal distribution, so that normality assumption seems to be reasonable, we can use repeated measures model here.

```
(b) stripchart(res ~ dat$A, method = 'stack')
```



These plots do not indicate any correlations of the error terms within a price level, and thus suggest that no interference effects are present.

(c) interaction.plot(dat\$S, dat\$A, dat\$Y)



We see that the rating curves for the stores do not appear to exhibit substantial departures from being parallel, hence, the assumption of no interactions appear to be reasonable.

```
(d)
     ab = dat1$S*dat1$A
     model1 = aov(Y~factor(A)+factor(S)+ab, data = dat1)
     anova(model1)
   ## Analysis of Variance Table
   ##
   ## Response: Y
   ##
                Df Sum Sq Mean Sq F value
                                              Pr(>F)
   ## factor(A)
                 2 67.48 33.740 52.9429 5.653e-07 ***
                 7 745.19 106.455 167.0410 1.056e-11 ***
   ## factor(S)
   ## ab
                 1
                     1.29
                            1.288
                                    2.0204
                                              0.1787
   ## Residuals 13
                     8.28
                            0.637
   ## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

```
qf(1-0.01, 1, 13)
## [1] 9.073806

pf(2.0204, 1, 13, lower.tail = FALSE)
## [1] 0.1787417
```

$$H_0:D=0$$
 VS. $H_1:D\neq 0$
$$F^* = \frac{SSAB^*/1}{SSrem*/(13)} = 1.288/0.637 = 2.0204$$

we can reject H_0 if $F^* > F(1 - 0.01; 1, 13) = 9.073806$, otherwise reject H_1 so that reject H_1 because $F^* < 9.073806$,

therefore, our conclusion implies that D equals zero, so there's no interactions, and P-value is 0.1787417.

```
model = aov(Y~ A + Error(S/A), data = dat)
 summary(model)
##
## Error: S
##
            Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 7 745.2 106.5
##
## Error: S:A
##
            Df Sum Sq Mean Sq F value Pr(>F)
            2 67.48 33.74 49.35 4.57e-07 ***
## A
## Residuals 14 9.57
                        0.68
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

```
(b) qf(1-0.05, 2, 14)

## [1] 3.738892

pf(49.35, 2, 14, lower.tail = FALSE)

## [1] 4.564874e-07
```

```
H_0:all \tau_j equal zero(j=1,2,3)
VS. H_1:not all \tau_j equal zero
F^* = \frac{MSA}{MSE} = 33.74/0.68 = 49.35
```

we can reject H_0 if $F^* > F(1 - 0.05; 2, 14) = 3.738892$, otherwise reject H_1 so that reject H_0 because $F^* > 3.738892$,

therefore, the mean sales of grape fruits differ for three price levels, and the P-value is $4.57\mathrm{e}\text{-}07$

```
(c) means = with(dat, by(Y, A, mean))
     D1 = means[1] - means[2]
     D2 = means[1] - means[3]
     D3 = means[2] - means[3]
     tukey = 1/sqrt(2)*qtukey(0.95, a, (a-1)*(r-1))
     tukey
   ## [1] 2.61728
     mstr.s = 0.68
     s = sqrt(2*mstr.s/(r))
   ## [1] 0.4123106
     c(D1-s*tukey, D1+s*tukey)
              1
   ## 0.7583676 2.9166324
     c(D2-s*tukey, D2+s*tukey)
             1
   ## 3.020868 5.179132
     c(D3-s*tukey, D3+s*tukey)
             2
   ## 1.183368 3.341632
```

$$\begin{split} \bar{Y}_{1\cdot\cdot\cdot} &= 55.4375, \bar{Y}_{2\cdot\cdot\cdot} = 53.6, \bar{Y}_{3\cdot\cdot\cdot} = 51.3375 \\ \hat{D}_1 &= \bar{Y}_{1\cdot\cdot\cdot} - \bar{Y}_{2\cdot\cdot\cdot} = 1.8375, \hat{D}_2 = \bar{Y}_{1\cdot\cdot\cdot} - \bar{Y}_{3\cdot\cdot\cdot} = 4.1, \hat{D}_3 = \bar{Y}_{2\cdot\cdot\cdot} - \bar{Y}_{3\cdot\cdot\cdot} = 2.2625 \end{split}$$

Because we estimate all pairwise comparisons, so we use tukey procedure

$$S = \sqrt{\frac{MSTR.S}{r}} * 2 = 0.4123106, Tukey = \frac{1}{\sqrt{2}} \text{qtukey} (1 - alpha, a, (a - 1) * (r - 1)) = 2.61728$$
 base on $\hat{D}_i \pm S * Tukey$
$$0.7583676 \le D_1 \le 2.9166324$$

$$3.020868 \le D_2 \le 5.179132$$

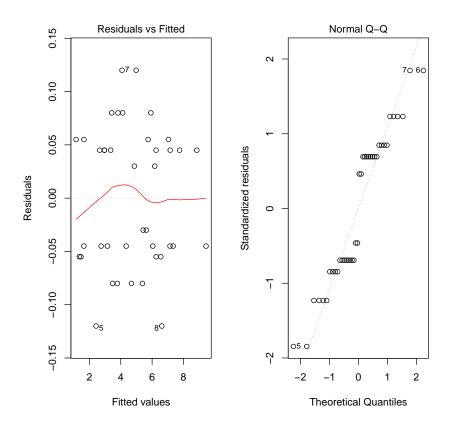
$$1.183368 \le D_3 \le 3.341632$$

(d)

$$\begin{split} \hat{E} &= \frac{S_r^2}{MSTR.S} \\ &= \frac{(s-1)MSS + s(r-1)*MSTR.S}{(sr-1)*MSTR.S} \\ &= 48.36189 \end{split}$$

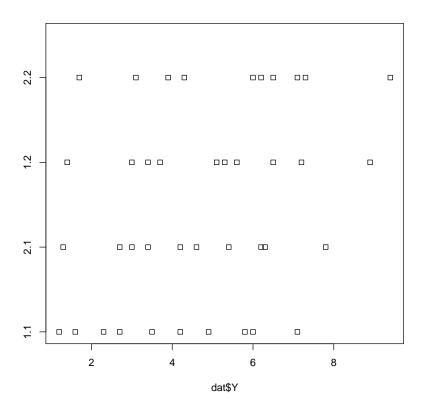
```
dat = read.table("CH27PR18.txt")
(a)
     names(dat) = c("Y", "S", "A", "B")
     dat$S = factor(dat$S)
     dat$A = factor(dat$A)
     dat$B = factor(dat$B)
     s = length(unique(dat$S))
     a = length(unique(dat$A))
     b = length(unique(dat$B))
     model = aov(Y^*S+A*B+A*S+B*S, data = dat)
     resid(model)
                 2
                        3
                             4
                                     5
                                            6
                                                   7
   ## -0.045 0.045 0.045 -0.045 -0.120 0.120 0.120 -0.120 0.080 -0.080
         11
             12
                    13 14
                                    15
                                           16
                                                  17
                                                         18
                                                                      20
   ## -0.080 0.080 -0.045 0.045 0.045 -0.045
                                              0.080 -0.080 -0.080
         21
                22
                       23
                              24
                                    25
                                           26
                                                  27
                                                         28
                                                               29
                                                                      30
       0.055 -0.055 -0.055 0.055 0.030 -0.030 -0.030
                                                      0.030 - 0.045
                                                                   0.045
                              34
             32
                   33
                                    35
                                           36
                                                         38
                                                  37
   ## 0.045 -0.045 0.055 -0.055 -0.055 0.055 -0.045 0.045 0.045 -0.045
```

```
par(mfrow = c(1,2))
plot(model, which = 1)
plot(model, which = 2)
```



The residuals versus fitted values plots shows no sign for unequal variance. And the QQ-plot indicates approximately normal distribution with slightly light tail, so that normality assumption seems to be reasonable, we can use model here.

```
(b) stripchart(dat$Y ~ dat$A*dat$B, method="stack")
```



These plots do not indicate any correlations of the error terms within AB, and thus suggest that no interference effects are present.

```
(a) model = aov(Y~ A*B+A*S+B*S+ Error(S/(A*B)), data = dat)
    summary(model)

##

## Error: S

## Df Sum Sq Mean Sq

## S 9 154.6 17.18

##

## Error: S:A

## Df Sum Sq Mean Sq

## A 1 3.025 3.0250
```

```
## A:S 9 2.035 0.2261

##

## Error: S:B

## Df Sum Sq Mean Sq

## B 1 11.449 11.449

## B:S 9 5.061 0.562

##

## Error: S:A:B

## Df Sum Sq Mean Sq F value Pr(>F)

## A:B 1 0.001 0.00100 0.053 0.823

## Residuals 9 0.169 0.01878
```

(b) From the plot, we know that treatment interaction effects are not present, but two treatment main effects are present.

```
(c) qf(1-0.005, 1, 9)
## [1] 13.61361

pf(0.053, 1, 9, lower.tail = FALSE)
## [1] 0.8230705
```

```
H_0:all (\alpha\beta)_{jk} equal zero
VS. H_1:not all (\alpha\beta)_{jk} equal zero
F^* = \frac{MSAB}{MSABS} = 0.001/0.01878 = 0.053
```

we can reject H_0 if $F^* > F(1 - 0.005; 1, 9) = 13.61361$, otherwise reject H_1 so that reject H_1 because $F^* < 13.61361$,

therefore, there's no two treatment interaction effect, and the P-value is 0.823

```
(d) qf(1-0.05, 1, 9)
## [1] 5.117355

pf(13.37904, 1, 9, lower.tail = FALSE)
## [1] 0.005253961
```

```
H_0: all \alpha_i equal zero(i=1,2)
VS. H_1:not all \alpha_i equal zero
F^* = \frac{MSA}{MSAS} = 3.0250/0.2261 = 13.37904
```

we can reject H_0 if $F^* > F(1 - 0.05; 1, 9) = 5.117355$,otherwise reject H_1 so that reject H_0 because $F^* > 5.117355$,

therefore, factor A main effect is present, and the P-value is 0.005253961

```
qf(1-0.05, 1, 9)
## [1] 5.117355

pf(20.37189, 1, 9, lower.tail = FALSE)
## [1] 0.001460304
```

 H_0 : all β_j equal zero(j=1,2) VS. H_1 :not all β_j equal zero $F^* = \frac{MSB}{MSBS} = 11.449/0.562 = 20.37189$

we can reject H_0 if $F^* > F(1 - 0.05; 1, 9) = 5.117355$,otherwise reject H_1 so that reject H_0 because $F^* > 5.117355$,

therefore, factor B main effect is present, and the P-value is 0.001460304

```
(e)    means = with(dat, by(Y, list(A,B), mean))
    L1 = means[2,1] - means[1,1]
    L2 = means[2,1] - means[1,2]
    L3 = means[2,2] - means[1,1]
    B = qt(1-0.05/(2*4), 9)
    B

## [1] 3.110935

msabs = 0.01878
    S = sqrt(2*msabs/(s))
    S

## [1] 0.06128621

c(L1-S*B, L1+S*B)
```

```
## [1] 0.3693426 0.7506574

c(L2-S*B, L2+S*B)

## [1] 0.8893426 1.2706574

c(L3-S*B, L3+S*B)

## [1] -0.7106574 -0.3293426

c(L4-S*B, L4+S*B)

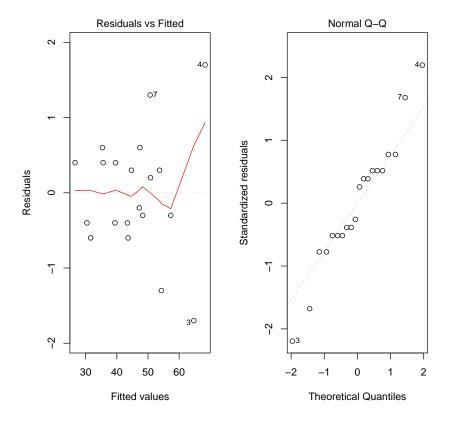
## [1] 1.429343 1.810657
```

$$\begin{split} \bar{Y}_{.11} &= 3.936, \bar{Y}_{.12} = 5.01, \\ \bar{Y}_{.21} &= 4.49, \bar{Y}_{.22} = 5.55 \\ \hat{L}_1 &= \bar{Y}_{.21} - \bar{Y}_{.11} = .56, \hat{L}_2 = \bar{Y}_{.12} - \bar{Y}_{.21} = 1.08, \\ \hat{L}_3 &= \bar{Y}_{.21} - \bar{Y}_{.12} = -.52, \hat{L}_4 = \bar{Y}_{.22} - \bar{Y}_{.11} = 1.62 \\ S &= \sqrt{\frac{MSABS}{s}} * 2 = 0.06128621, B = t(1 - \alpha/(2*4), (a-1)(b-1)(s-1)) = 3.110935 \\ \text{base on} \hat{L}_i \pm S*B \\ 0.3693426 \leq L_1 \leq 0.7506574 \\ 0.8893426 \leq L_2 \leq 1.2706574 \\ -0.7106574 \leq L_3 \leq -0.3293426 \\ 1.429343 \leq L_4 \leq 1.810657 \end{split}$$

The treatment of high dose of both drugs has the most significant reduction in pain intensity compared to the treatments of only one drug has high dose, and a significant difference exists in the mean effects of two drugs used alone.

```
(a) dat = read.table("CH27PR20.txt")
    names(dat) = c("Y", "S", "A", "B")
    dat$S = factor(dat$S)
    dat$A = factor(dat$A)
    dat$B = factor(dat$B)
```

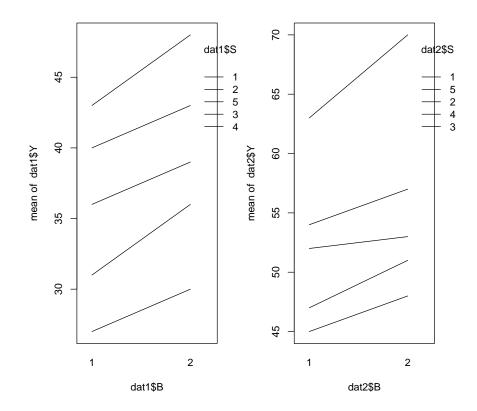
```
s = length(unique(dat$S))
  a = length(unique(dat$A))
  b = length(unique(dat$B))
  model = aov(Y^{\sim} A*B+(A/S), data = dat)
  resid(model)
      1
                            5
                                      7
                                            8
                 3
                      4
                                 6
                                                 9
                                                      10
                                                           11
                                                                12
                                                                      13
                                                                           14
##
   -0.6
             -1.7
                    1.7
                         0.4 - 0.4
                                    1.3 -1.3 -0.6
                                                    0.6
                                                          0.3 -0.3
                                                                     0.4 - 0.4 - 0.2
##
     16
                18
                     19
                           20
           17
    0.2
         0.4 - 0.4
                    0.3 -0.3
  par(mfrow = c(1,2))
  plot(model, which = 1)
  plot(model, which = 2)
```



The residuals versus fitted values plots shows no sign for unequal variance. And the QQ-plot indicates approximately normal distribution with

slightly heavy tail, so that normality assumption seems to be reasonable, we can use model here.

```
(b) dat1 = dat[ which(dat$A == 1), ]
  dat2 = dat[ which(dat$A == 2), ]
  par( mfrow = c(1, 2))
  interaction.plot(dat1$B, dat1$S, dat1$Y, lty = 1)
  interaction.plot(dat2$B, dat2$S, dat2$Y, lty = 1)
```



There's no evidence any interactions between field and treatments, so split-plot design can be used here.

```
(a) model = aov(Y~ A*B+Error(A/S), data = dat)
summary(model)
```

```
##
## Error: A
## Df Sum Sq Mean Sq
## A 1 1394 1394
##
## Error: A:S
##
           Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 8 837.6 104.7
##
## Error: Within
                                   Pr(>F)
##
      Df Sum Sq Mean Sq F value
           1 68.45 68.45 45.633 0.000144 ***
## A:B 1 0.05 0.05 0.033 0.859674
## Residuals 8 12.00
                     1.50
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

(b) From the plot, we know that treatment interaction effects are not present, but two treatment main effects are present.

```
(c) qf(1-0.05, 1, 8)

## [1] 5.317655

pf(0.033, 1, 8, lower.tail = FALSE)

## [1] 0.8603687
```

$$H_0$$
:all $(\alpha\beta)_{jk}$ equal zero VS. H_1 :not all $(\alpha\beta)_{jk}$ equal zero $F^* = \frac{MSAB}{MSB.W(A)} = 0.05/1.5 = 0.033$

we can reject H_0 if $F^* > F(1 - 0.05; 1, 8) = 5.317655$,otherwise reject H_1 so that reject H_1 because $F^* < 5.317655$,

therefore, there's no two treatment interaction effect, and the P-value is 0.8603687

```
(d) qf(1-0.05, 1, 8)
## [1] 5.317655
pf(13.31423, 1, 8, lower.tail = FALSE)
## [1] 0.006505011
```

 H_0 : all α_i equal zero(i=1,2) VS. H_1 :not all α_i equal zero $F^* = \frac{MSA}{MSW(A)} = 1394/104.7 = 13.31423$

we can reject H_0 if $F^* > F(1 - 0.05; 1, 8) = 5.317655$,otherwise reject H_1 so that reject H_0 because $F^* > 5.317655$,

therefore, factor A effect is present, and the P-value is 0.00650501

```
qf(1-0.05, 1, 8)
## [1] 5.317655

pf(45.63333, 1, 8, lower.tail = FALSE)
## [1] 0.0001442759
```

 H_0 : all β_j equal zero(j=1,2) VS. H_1 :not all β_j equal zero $F^* = \frac{MSB}{MSB.W(A)} = 68.45/1.5 = 45.63333$

we can reject H_0 if $F^* > F(1-0.05;1,8) = 5.317655$, otherwise reject H_1 so that reject H_0 because $F^* > 5.317655$,

therefore, factor B main effect is present, and the P-value is 0.0001442759

```
(e)    means1 = with(dat, by(Y, A , mean))
    means2 = with(dat, by(Y, B , mean))
    L1 = means1[1] - means1[2]
    L2 = means2[1] - means2[2]
    B = qt(1-0.1/(2*2), 8)
    B

## [1] 2.306004

msb.wa = 1.5
mswa = 104.7
S1 = sqrt(2*mswa/(b*s))
S2 = sqrt(2*msb.wa/(a*s))
S1

## [1] 4.576024
```

```
## [1] 0.5477226

c(L1-S1*B, L1+S1*B)

## 1 1

## -27.252331 -6.147669

c(L2-S2*B, L2+S2*B)

## 1 1

## -4.96305 -2.43695
```

$$\bar{Y}_{.1} = 37.3, \bar{Y}_{.2} = 54,$$

$$\bar{Y}_{.1} = 43.8, \bar{Y}_{.2} = 47.5$$

$$\hat{L}_1 = \bar{Y}_{.1} - \bar{Y}_{.2} = -16.7, \hat{L}_2 = \bar{Y}_{.1} - \bar{Y}_{.2} = -3.7$$

$$B = t(1 - \alpha/(2*2), a(b-1)(s-1)) = 2.306004$$

$$S1 = \sqrt{\frac{MSW(A)}{bs}} * 2 = 4.576024, S2 = \sqrt{\frac{MSB.W(A)}{as}} * 2 = 0.5477226$$

$$\text{base on } \hat{L}_i \pm S*B$$

$$-27.252331 \le L_1 \le -6.147669$$

$$-4.96305 \le L_2 \le -2.43695$$

Irrigation method 2 is significantly better than Irrigation method 1, and fertilizer 2 is significantly better than fertilizer 1.

7 27.22

$$\begin{split} SSTO &= \sum_{i} \sum_{j} (Y_{ij} - \bar{Y}_{..})^{2} \\ &= \sum_{i} \sum_{j} (Y_{ij} - \bar{Y}_{i.} + \bar{Y}_{i.} - \bar{Y}_{..})^{2} \\ &= \sum_{i} \sum_{j} (Y_{ij} - \bar{Y}_{i.})^{2} + \sum_{i} \sum_{j} (\bar{Y}_{i.} - \bar{Y}_{..})^{2} + 2 \sum_{i} \sum_{j} (Y_{ij} - \bar{Y}_{i.})(\bar{Y}_{i.} - \bar{Y}_{..}) \\ &= \sum_{i} \sum_{j} (Y_{ij} - \bar{Y}_{i.})^{2} + \sum_{i} \sum_{j} (\bar{Y}_{i.} - \bar{Y}_{..})^{2} + 2 \sum_{i} [(\bar{Y}_{i.} - \bar{Y}_{..}) * \sum_{j} (Y_{ij} - \bar{Y}_{i.})] \\ &(\text{since } \sum_{j} (Y_{ij} - \bar{Y}_{i.}) = 0) \\ &= \sum_{i} \sum_{j} (Y_{ij} - \bar{Y}_{i.})^{2} + r \sum_{i} (\bar{Y}_{i.} - \bar{Y}_{..})^{2} \\ &= SSS + SSW \end{split}$$

8 Extra Problem

Base on:

$$\begin{split} Y_{ij} &= \mu... + \rho_i + \tau_j + \epsilon_{ij} \\ \bar{Y}_{i\cdot} &= \mu... + \rho_i + \bar{\epsilon}_{i\cdot} \\ \bar{Y}_{\cdot j} &= \mu... + \bar{\rho}_{\cdot} + \tau_j + \bar{\epsilon}_{\cdot j} \\ \bar{Y}_{\cdot \cdot j} &= \mu... + \bar{\rho}_{\cdot} + \bar{\epsilon}_{\cdot \cdot} \end{split}$$

So that:

$$\begin{split} E(SSS) &= r \sum_{i} (E(\bar{Y}_{i\cdot} - \bar{Y}_{\cdot\cdot})^2) \\ &= r \sum_{i} (Var(\bar{Y}_{i\cdot} - \bar{Y}_{\cdot\cdot}) + [E(\bar{Y}_{i\cdot} - \bar{Y}_{\cdot\cdot})]^2) \\ &(\text{since } E(\bar{Y}_{i\cdot} - \bar{Y}_{\cdot\cdot}) = 0) \\ &= r \sum_{i} (Var(\bar{Y}_{i\cdot}) + Var(\bar{Y}_{\cdot\cdot}) - 2Cov((\bar{Y}_{i\cdot}, \bar{Y}_{\cdot\cdot})) \\ &= r \sum_{i} ((\sigma_{\rho}^2 + \frac{\sigma^2}{r}) + (\frac{\sigma_{\rho}^2}{s} + \frac{\sigma^2}{rs}) - 2 * (Cov(\rho_i, \bar{\rho}_{\cdot}) + Cov(\bar{\epsilon}_{i\cdot}, \bar{\rho}_{\cdot}) + Cov(\rho_i, \bar{\epsilon}_{\cdot\cdot}) + Cov(\bar{\epsilon}_{i\cdot}, \bar{\epsilon}_{\cdot\cdot})) \\ &= rs * ((\sigma_{\rho}^2 + \frac{\sigma^2}{r}) + (\frac{\sigma_{\rho}^2}{s} + \frac{\sigma^2}{rs}) - 2 * (\frac{\sigma_{\rho}^2}{s} + 0 + 0 + \frac{\sigma^2}{rs})) \\ &= (s - 1)\sigma^2 + r(s - 1)\sigma_{\rho}^2 \\ E(MSS) &= \frac{E(SSS)}{s - 1} \\ &= \sigma^2 + r\sigma_{\rho}^2 \end{split}$$

Likewise:

$$\begin{split} E(SSTR) &= s \sum_{j} (E(\bar{Y}_{\cdot j} - \bar{Y}_{\cdot \cdot})^{2}) \\ &= s \sum_{j} (Var(\bar{Y}_{\cdot j} - \bar{Y}_{\cdot \cdot}) + [E(\bar{Y}_{\cdot j} - \bar{Y}_{\cdot \cdot})]^{2}) \\ &= s \sum_{j} (Var(\bar{Y}_{\cdot j}) + Var(\bar{Y}_{\cdot \cdot}) - 2Cov((\bar{Y}_{\cdot j}, \bar{Y}_{\cdot \cdot}) + \tau_{j}) \\ &= s \sum_{j} ((\frac{\sigma_{\rho}^{2}}{s} + \frac{\sigma^{2}}{s}) + (\frac{\sigma_{\rho}^{2}}{s} + \frac{\sigma^{2}}{rs}) - 2 * (\frac{\sigma_{\rho}^{2}}{s} + 0 + 0 + \frac{\sigma^{2}}{rs}) + \tau_{j}) \\ &= (r - 1)\sigma^{2} + s \sum_{j} \tau_{j}^{2} \\ E(MSTR) &= \frac{E(SSTR)}{r - 1} \\ &= \sigma^{2} + s \frac{\sum_{j} \tau_{j}^{2}}{r - 1} \end{split}$$

$$E(SSTR.S) = \sum_{i} \sum_{j} (E(Y_{ij} - \bar{Y}_{i.} - \bar{Y}_{.j} + \bar{Y}_{..})^{2})$$

$$= \sum_{i} \sum_{j} (Var(Y_{ij} - \bar{Y}_{i.} - \bar{Y}_{.j} + \bar{Y}_{..}) + [E(Y_{ij} - \bar{Y}_{i.} - \bar{Y}_{.j} + \bar{Y}_{..})]^{2})$$

$$= \sum_{i} \sum_{j} (Var(Y_{ij} - \bar{Y}_{i.} - \bar{Y}_{.j} + \bar{Y}_{..}))$$

$$= (r - 1)(s - 1)\sigma^{2}$$

$$E(MSTR.S) = \frac{E(SSTR.S)}{(r - 1)(s - 1)}$$

$$= \sigma^{2}$$