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#### Untitled

#### 2024-03-28

```
level <- c(rep(1,5), rep(2,5), rep(3,5))
a <- c(49,73,58,38,42,31,40,43,44,20,46,41,58,31,65)
table <- cbind(level, a)
table <- data.frame(table)
```

## 분산분석표

```
aov1 <- aov(a~level)
summary(aov1)
```

```
## Df Sum Sq Mean Sq F value Pr(>F)
## level 1 36.1 36.1 0.179 0.679
## Residuals 13 2622.8 201.8
```

```
out1 = Im(a~con1+con2, data = table)
summary(out1)
```

```
##
## Im(formula = a \sim con1 + con2, data = table)
##
## Residuals:
##
   Min
         1Q Median
                      3Q
                              Max
  -17.2 -8.6 -2.2
                        7.9 21.0
##
## Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 45.267 3.268 13.853 9.62e-09 ***
               -1.900
                          4.002 - 0.475
## con1
                                         0.6435
## con2
               -4.833
                         2.311 -2.092 0.0584 .
## ---
## Signif. codes: 0 '*** 0.001 '** 0.05 '.' 0.1 ' 1
## Residual standard error: 12.66 on 12 degrees of freedom
## Multiple R-squared: 0.2772, Adjusted R-squared: 0.1567
## F-statistic: 2.301 on 2 and 12 DF, p-value: 0.1427
```

# 중회귀분석의 분산분석표

```
out2 <- aov(out1)
summary(out2)</pre>
```

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검정결과가 동일하다.

## 순서

```
out3 = Im(a~con2+con1, data = table)
out4 <- aov(out3)
summary(out4)</pre>
```

순서에 상관이 없이 동일하다.