

# DSA 8070 R Session 7: Repeated Measures Analysis

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## Read the data

```
dat <- read.table("dog1.txt")
temp <- array(dim = c(144, 4))
temp[, 1] <- rep(dat$V1, 4)
temp[, 2] <- rep(dat$V2, 4)
temp[, 3] <- rep(c(1, 5, 9, 13), each = 36)
temp[, 4] <- c(dat$V3, dat$V4, dat$V5, dat$V6)
dat2 <- data.frame(temp)
names(dat2) <- c("Treatment", "Dog_id", "Time", "Response")
dat2$Treatment <- as.factor(dat2$Treatment)
dat2$Dog_id <- as.factor(dat2$Dog_id)
dat2$Time <- as.factor(dat2$Time)
```

## Split-plot ANOVA

```
# computing the cell means (by treatment and time combinations)
tapply(dat2$Response, list(dat2$Treatment, dat2$Time), mean)
```

```
##           1           5           9          13
## 1 4.111111 4.400000 5.066667 4.722222
## 2 3.600000 4.200000 4.500000 4.225000
## 3 3.644444 4.011111 3.977778 4.044444
## 4 3.540000 3.620000 3.660000 3.460000
```

```

# interaction plot
par(las = 1, mgp = c(2.2, 1, 0), mar = c(3.6, 3.6, 0.8, 0.6))
with(dat2, interaction.plot(x.factor = Time, trace.factor = Treatment,
                           response = Response, col = 1:4, lwd = 1.5))

# Split-plot anova
library(lmerTest)

```

```
## Loading required package: lme4
```

```
## Loading required package: Matrix
```

```
## Warning: replacing previous import 'lifecycle::last_warnings' by
## 'rlang::last_warnings' when loading 'tibble'
```

```
## Warning: replacing previous import 'lifecycle::last_warnings' by
## 'rlang::last_warnings' when loading 'pillar'
```

```
##
```

```
## Attaching package: 'lmerTest'
```

```
## The following object is masked from 'package:lme4':
```

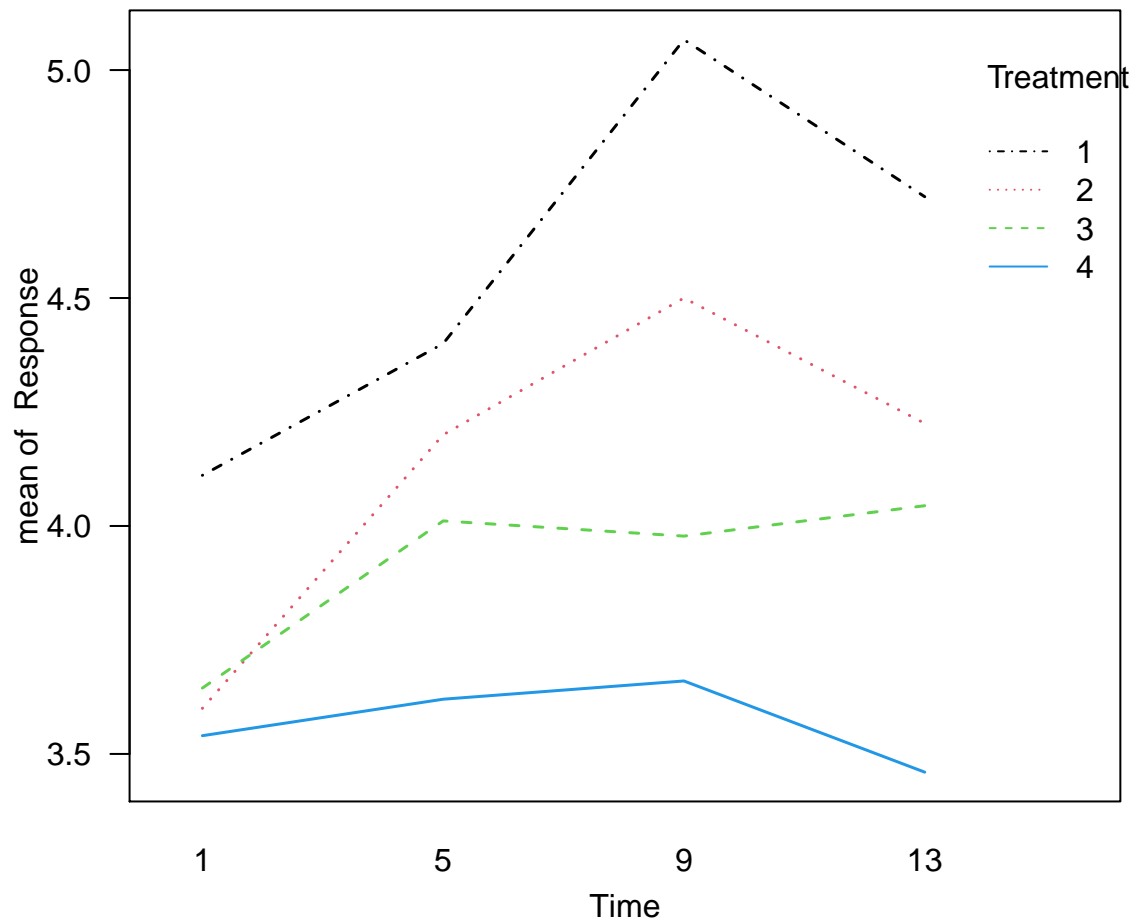
```
##
```

```
##      lmer
```

```
## The following object is masked from 'package:stats':
```

```
##
```

```
##      step
```



```
fit <- lmer(Response ~ Treatment * Time + (1 | Dog_id), data = dat2)
anova(fit)
```

```
## Type III Analysis of Variance Table with Satterthwaite's method
##               Sum Sq Mean Sq NumDF DenDF F value    Pr(>F)
## Treatment      3.3396  1.11319     3    32   6.0038 0.002297 **
## Time           6.2043  2.06811     3    96  11.1540 2.404e-06 ***
## Treatment:Time  3.4397  0.38219     9    96   2.0613 0.040573 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

## MANOVA

```
dat <- read.table("dog1.txt")
out <- manova(cbind(V3, V4, V5, V6) ~ as.factor(V1), data = dat)
summary(out, test = "Wilks")
```

```
##               Df  Wilks approx F num Df den Df  Pr(>F)
## as.factor(V1)  3 0.48452    2.022    12 77.018 0.03316 *
## Residuals      32
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

## Mixed Model with AR(1) temporal correlation structure

```
library(nlme)

##
## Attaching package: 'nlme'

## The following object is masked from 'package:lme4':
##
##      lmList

fit1 = gls(Response ~ Treatment * Time,
            correlation = corCompSymm(form = ~ 1 | Dog_id), data = dat2)
fit2 = gls(Response ~ Treatment * Time,
            correlation = corAR1(form = ~ 1 | Dog_id), data = dat2)
anova(fit1, fit2)

##      Model df      AIC      BIC    logLik
## fit1      1 18 275.8063 327.1429 -119.9032
## fit2      2 18 277.5811 328.9177 -120.7906
```