# 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$Dog_id <- as.factor(dat2$Treatment)
dat2$Time <- as.factor(dat2$Time)</pre>
```

## 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 4.7222222

## 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

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))

library(lmerTest)

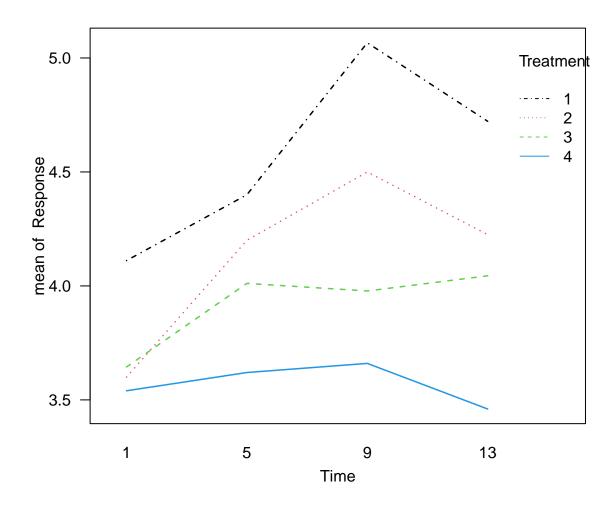
## Loading required package: lme4

## Loading required package: Matrix

## ## 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)</pre>
```

```
## 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 '**' 0.05 '.' 0.1 ' ' 1
```

### **MANOVA**

## fit2

```
summary(out, test = "Wilks")
##
                    Wilks approx F num Df den Df Pr(>F)
## as.factor(V1) 3 0.48452
                            2.022
                                     12 77.018 0.03316 *
## Residuals
## ---
## 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
```

out <- manova(cbind(V3, V4, V5, V6) ~ as.factor(V1), data = dat)</pre>

2 18 277.5811 328.9177 -120.7906