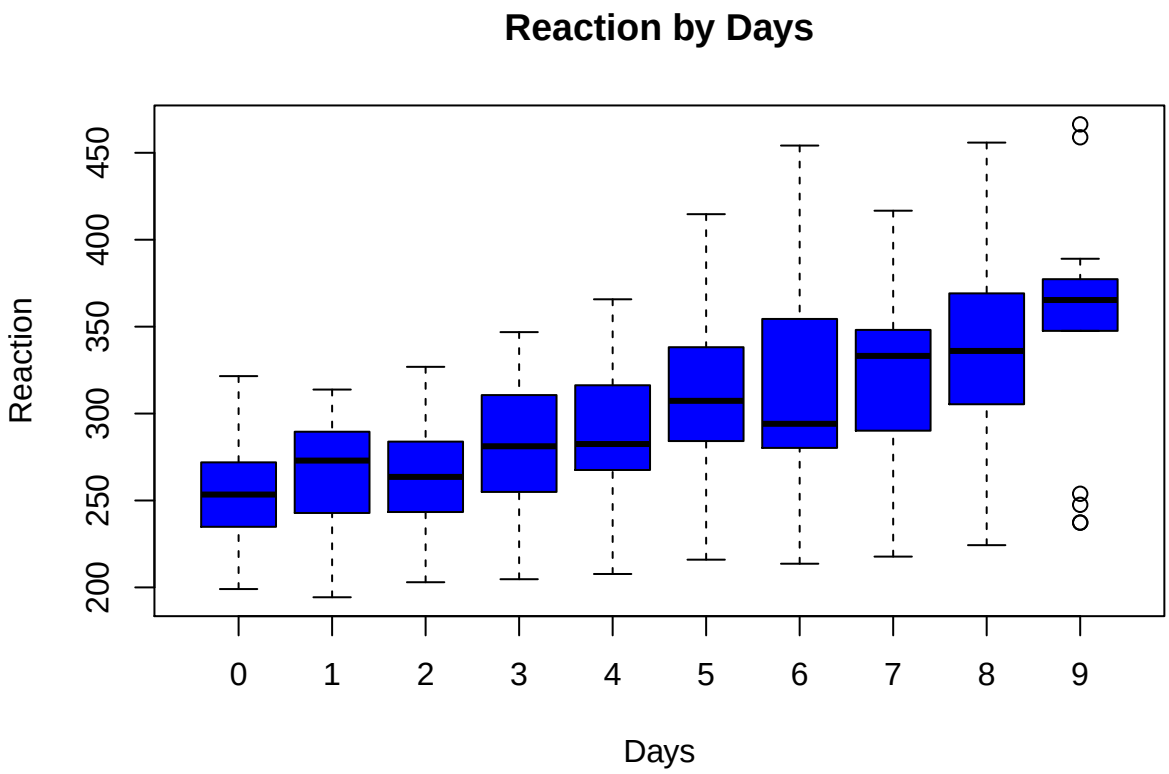
## P values from linear and linear mixed models

As its name implies, P values are provided by the `lmerTest` package, as will be illustrated with the documentation example here.

```
require(lmerTest)
boxplot(Reaction~Days, data=sleepstudy, main="Reaction by Days",
        xlab="Days", ylab="Reaction", col="blue", border="black")
```




We see a trend of `Reaction` by `Days`, so it is reasonable to fit a simple linear regression,

```
l <- lm(Reaction ~ Days, sleepstudy)
s <- summary(l)
s
```

```
##
## Call:
## lm(formula = Reaction ~ Days, data = sleepstudy)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -110.848  -27.483    1.546   26.142  139.953
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept)  251.405      6.610  38.033  < 2e-16 ***
## Days          10.467      1.238   8.454 9.89e-15 ***
## ---
```

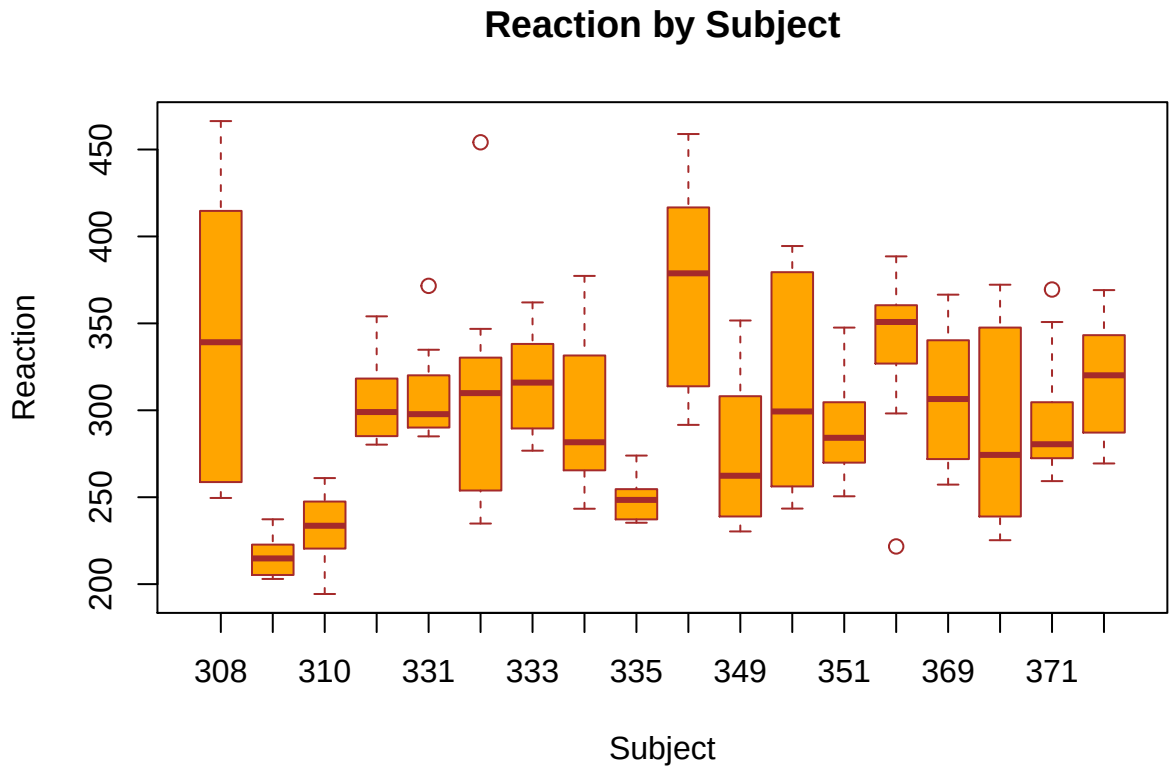
```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 47.71 on 178 degrees of freedom
## Multiple R-squared:  0.2865, Adjusted R-squared:  0.2825
## F-statistic: 71.46 on 1 and 178 DF,  p-value: 9.894e-15
```

```
names(s)
```

```
##  [1] "call"          "terms"         "residuals"     "coefficients"
##  [5] "aliased"       "sigma"         "df"            "r.squared"
##  [9] "adj.r.squared" "fstatistic"    "cov.unscaled"
```

Consider now the `Subject` effect. From

```
boxplot(Reaction~Subject, data=sleepstudy, main="Reaction by Subject",
        xlab="Subject", ylab="Reaction", col="orange", border="brown")
```




it is more approriate to fit a random effect model

```
f <- lme4::lmer(Reaction ~ Days + (Days | Subject), sleepstudy)
s <- summary(f)
s
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: Reaction ~ Days + (Days | Subject)
##    Data: sleepstudy
##
## REML criterion at convergence: 1743.6
##
## Scaled residuals:
```

```
##     Min      1Q  Median      3Q     Max
## -3.9536 -0.4634  0.0231  0.4633  5.1793
##
## Random effects:
##  Groups   Name        Variance Std.Dev. Corr
##  Subject  (Intercept) 611.90   24.737
##           Days         35.08    5.923   0.07
##  Residual             654.94   25.592
## Number of obs: 180, groups:  Subject, 18
##
## Fixed effects:
##             Estimate Std. Error t value
## (Intercept)  251.405      6.824  36.843
## Days          10.467      1.546   6.771
##
## Correlation of Fixed Effects:
##      (Intr)
## Days -0.138
```

```
names(s)
```

```
##  [1] "methTitle"    "objClass"     "devcomp"      "isLmer"
##  [5] "useScale"     "logLik"       "family"       "link"
##  [9] "ngrps"        "coefficients" "sigma"        "vcov"
## [13] "varcor"       "AICtab"       "call"         "residuals"
## [17] "fitMsgs"      "optinfo"
```

We saw the same estimate of effect but a larger standard error for `Days` in the linear mixed model compared to that in the linear regression model. We then use `lmer` from `lmerTest`.

```
m <- lmerTest::lmer(Reaction ~ Days + (Days | Subject), sleepstudy)
class(m)
```

```
## [1] "lmerModLmerTest"
## attr(,"package")
## [1] "lmerTest"
```

```
s <-summary(m)
names(s)
```

```
##  [1] "methTitle"    "objClass"     "devcomp"      "isLmer"
##  [5] "useScale"     "logLik"       "family"       "link"
##  [9] "ngrps"        "coefficients" "sigma"        "vcov"
## [13] "varcor"       "AICtab"       "call"         "residuals"
## [17] "fitMsgs"      "optinfo"
```

```
with(s,coefficients)[2,5]
```

```
## [1] 3.273014e-06
```

Compared to its counterpart in `lme4`, `lmerTest` provides P value.