## Model results and diagnostics

There are many functions available to produce predicted values and diagnostics. For additional commands not listedhere, see help (influence. measures)).

### **Predicted values**

```
mod1 = lm(y ~ x, data=ds)
predicted.varname = predict(mod1)
```

### **Residuals**

```
mod1 = lm(y ~ x, data=ds)
residual.varname = residuals(mod1)
```

### Standardized and Studentized residuals

```
mod1 = lm(y ~ x, data=ds)
standardized.resid.varname = rstandard(mod1)
studentized.resid.varname = rstudent(mod1)
```

### Leverage

```
mod1 = lm(y ~ x, data=ds)
leverage.varname = hatvalues(mod1)
```

#### Cook's distance

```
mod1 = lm(y ~ x, data=ds)
cookd.varname = cooks.distance(mod1)
```

### **DFFITs**

```
mod1 = lm(y ~ x, data=ds)
dffits.varname = dffits(mod1)
```

# **Diagnostic plots**

```
mod1 = lm(y ~ x, data=ds)
par(mfrow=c(2,2)) # display 2 x 2 matrix of graphs
plot(mod1)
```

# **Heteroscedasticity tests**

```
library(lmtest)
bptest(y ~ x1 + ... + xk, data=ds)
```

#### **Parameter estimates**

```
mod1 = lm(y ~ x, data=ds)
coeff.mod1 = coef(mod1)
```

# Standardized regression coefficients

```
library(QuantPsyc)
mod1 = lm(y ~ x)
lm.beta(mod1)
```

## Confidence interval for parameter estimates

```
mod1 = lm(y ~ x, data=ds)
confint(mod1)
```

### **Prediction limits**

```
mod1 = lm(y ~ ..., data=ds)
pred.w.lowlim = predict(mod1, interval="prediction")[,2]
```

## Covariance matrix of parameter estimates

```
mod1 = lm(y ~ x, data=ds)
vcov(mod1)
```

# **Correlation matrix of parameter estimates**

```
mod1 = lm(y ~ x, data=ds)
mod1.cov = vcov(mod1)
mod1.cor = cov2cor(mod1.cov)
```