

One-Way ANOVA Diagnostics

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Assumptions of a One-Way ANOVA

1. Independence of observations
2. Variances (SD) between groups are the same
3. (F-test) Normality of responses (or errors)

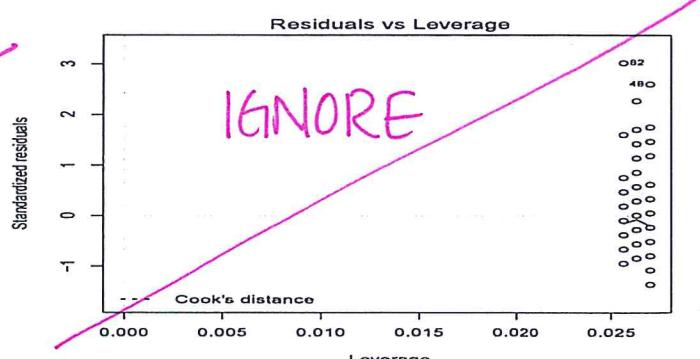
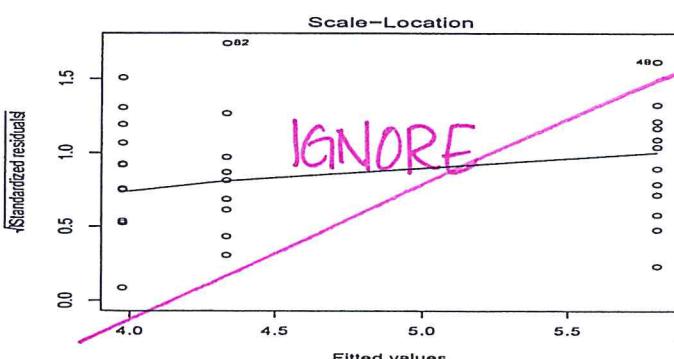
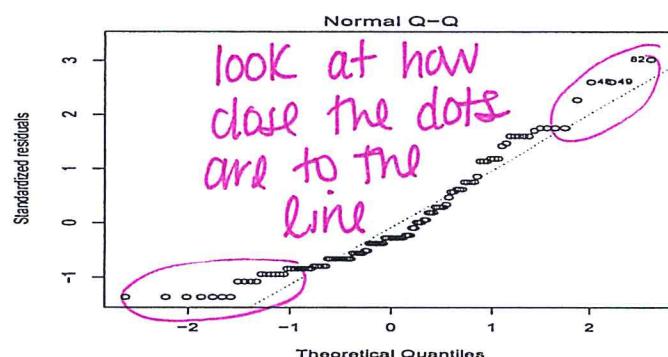
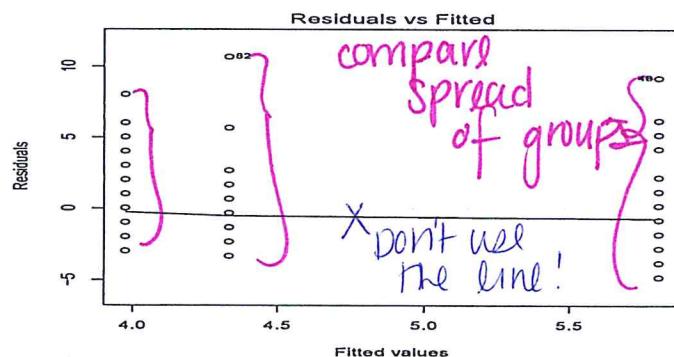
Balanced Design:

Every group has the same sample size.

* Not an assumption, but can influence your test results!

R's built-in diagnostic plots:

```
require(heplots)
data(MockJury)
mockJury.model <- lm(Years ~ Attr, data = MockJury)
par(mfrow = c(2, 2))
plot(mockJury.model)
```



- Residuals vs Fitted Plot

Displays residuals on y-axis : fitted values on x-axis. Look for: equal spread of dots between groups!

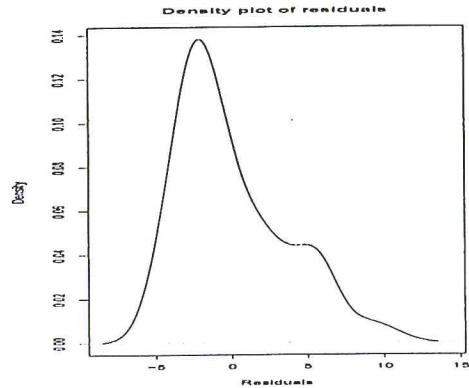
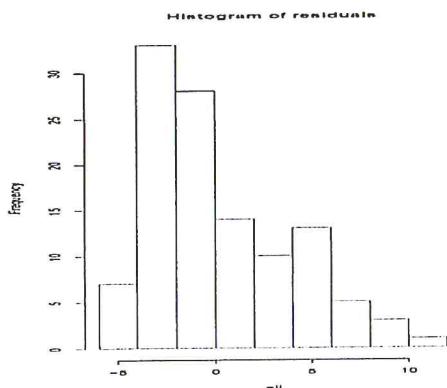
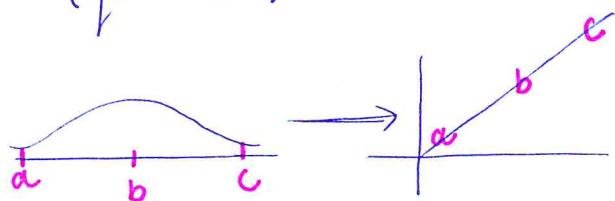
- Normal Q-Q Plot

Displays errors (residuals) on a Normal distribution (quantiles)

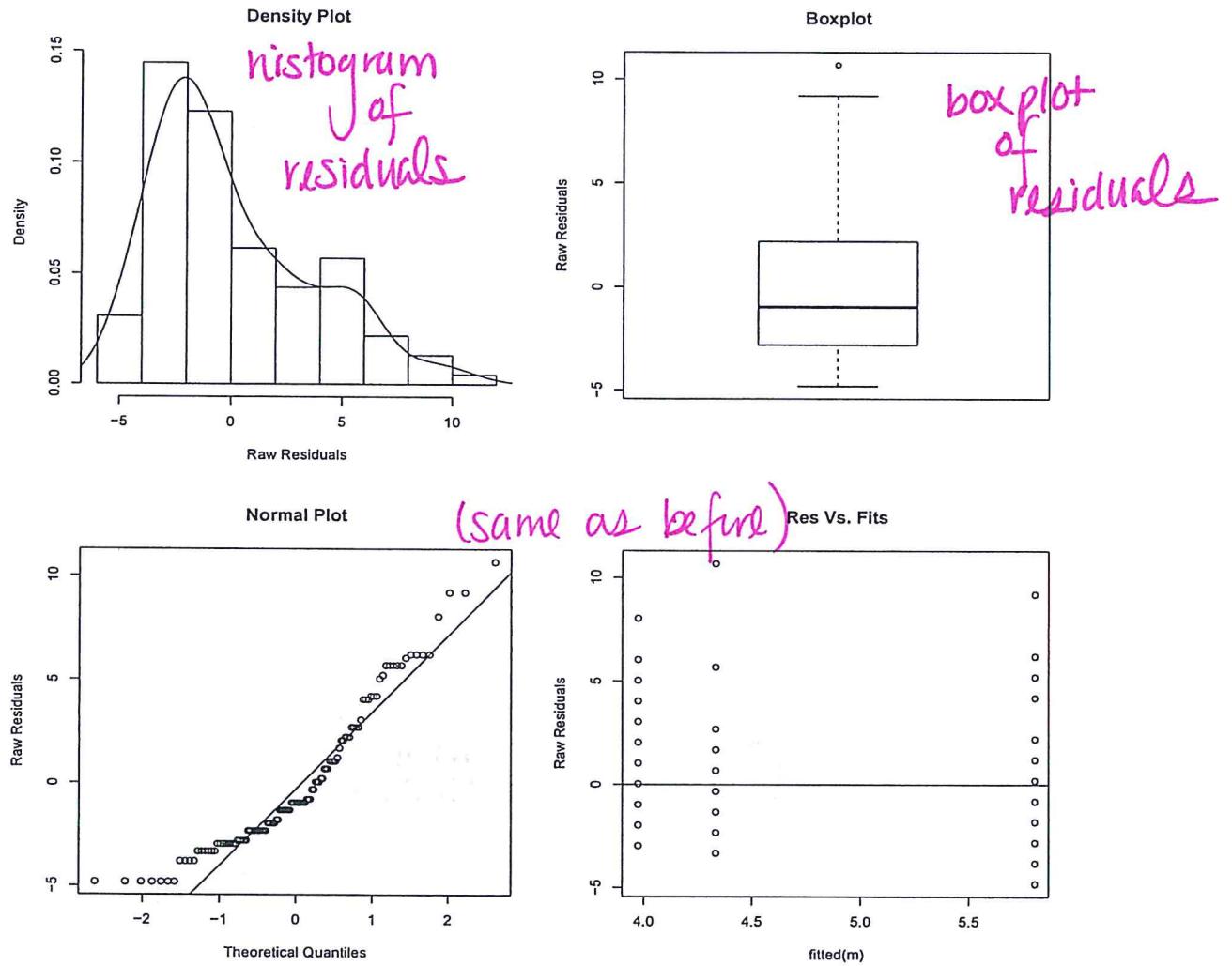
Look for: points following the line fairly close

Another way to look at the distribution of the residuals:

```
par(mfrow = c(1, 2))
eij <- residuals(mockJury.model)
hist(eij, main = "Histogram of residuals")
plot(density(eij), main = "Density plot of residuals", ylab = "Density",
      xlab = "Residuals")
```



Al's diagANOVA Function



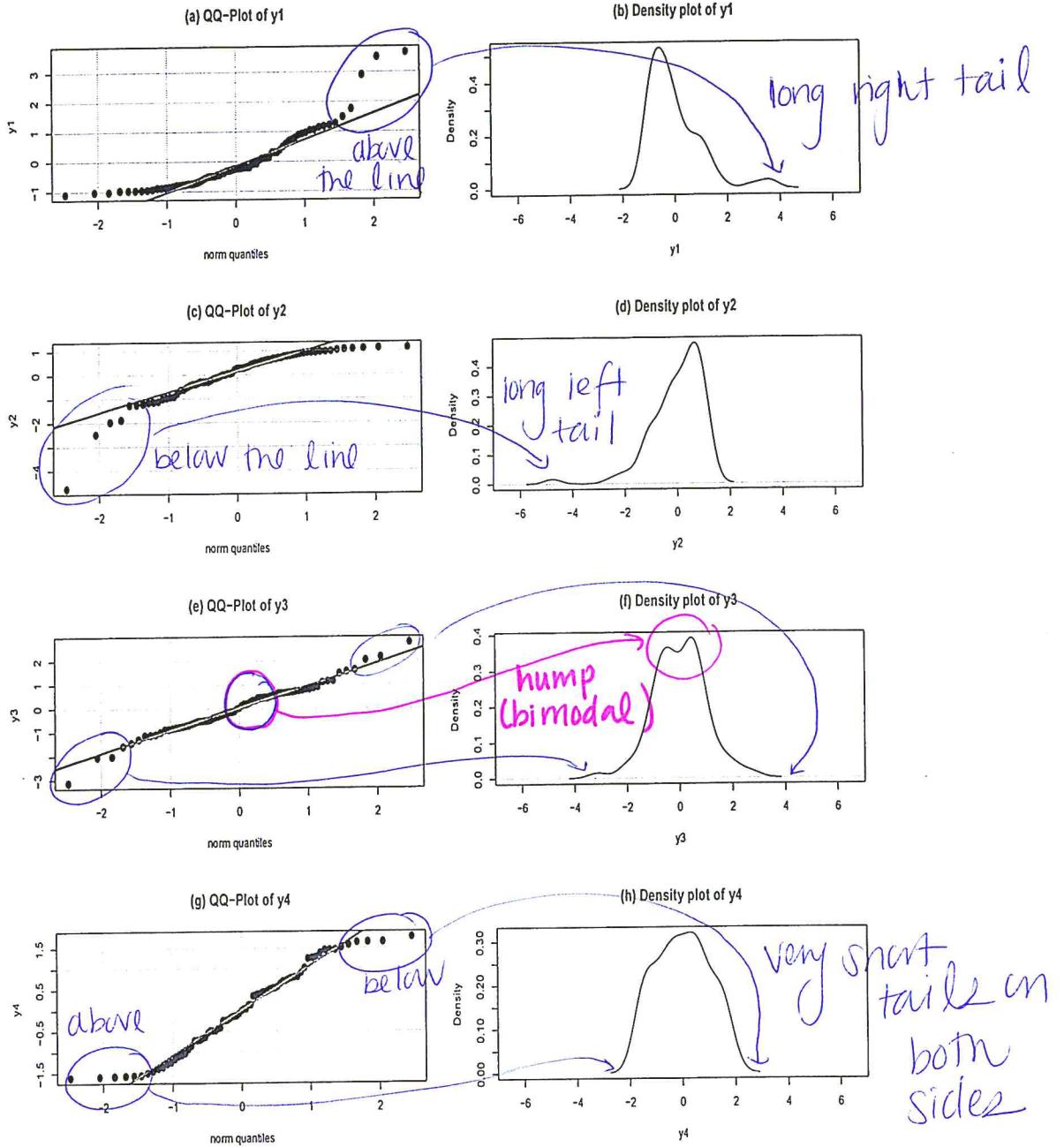
```
## [1] "In this sample of size n=114, correlation of the residuals in the qq-plot is r=0.957673"
## [1] "In the following table, if r < critical.r, then the qq-plot suggests the residuals are not normal:"
##   n  critical.r
## 1 5    0.832
## 2 10   0.880
## 3 15   0.911
## 4 20   0.929
## 5 25   0.941
## 6 30   0.949
## 7 40   0.960
## 8 50   0.966
## 9 60   0.971
## 10 75  0.976
##
## Shapiro-Wilk normality test
## 
## data: r
## W = 0.9, p-value = 2e-06
```

correlation check for Normality

If you use diagANOVA with do.tests = TRUE you get these!

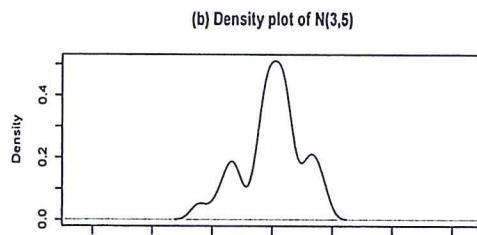
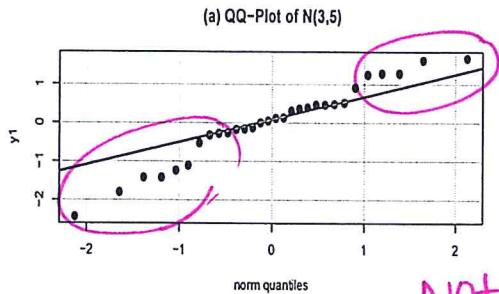
] you won't use this.

Some examples of Q-Q plots:

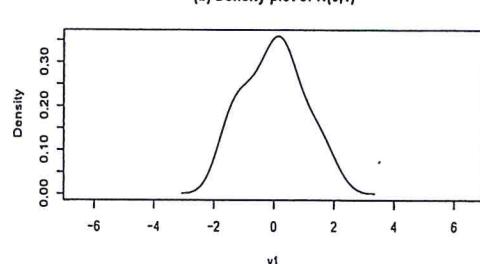
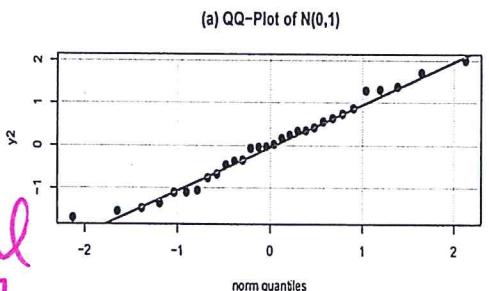


Some examples of Normal Q-Q plots:

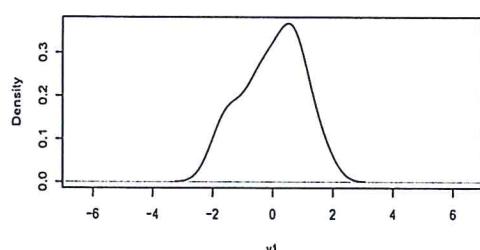
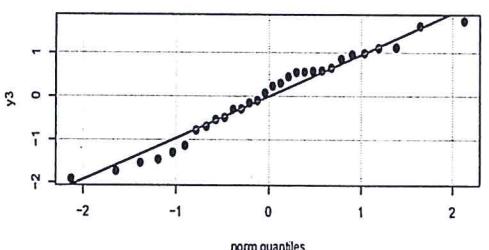
These are Q-Q plots that come from Normal data!



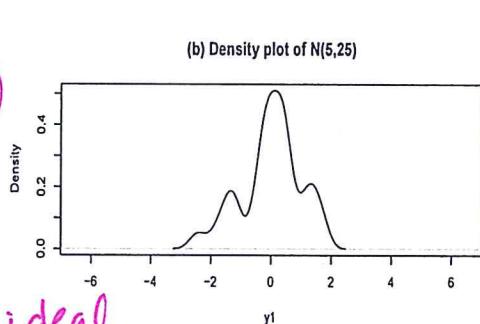
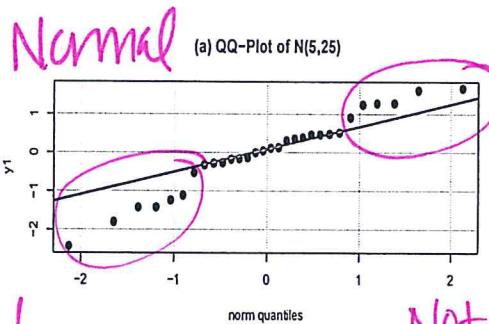
NOT ideal, but these data are Normal!



(a) QQ-Plot of $N(5,10)$



Keep in mind even



Not ideal, but these data are Normal!

Checking the balance of the design:

`table(MockJury$Attr)` ← table will get you counts in each group.

```
##          Beautiful      Average      Unattractive
##            39                  38                  37
```

Unequal sample sizes = unbalanced design

