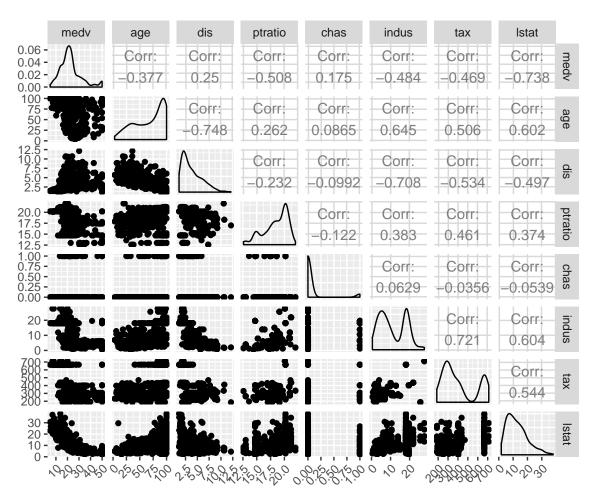
Notes: MS 204 Chapter 6.2

Overview

- Statistical significance
- Model selection
- Model validation

Multiple linear regression: Boston city home prices

```
Ex: X_1 = \text{crim}, X_2 = rm, \dots, Y = \text{medv}
library(mosaic); library(tidyverse); library(MASS)
data (Boston)
dim(Boston)
## [1] 506 14
Boston.reg <- Boston %>%
  dplyr::select(medv, age, dis, ptratio, chas, indus, tax, lstat)
head(Boston.reg)
     medv age
                  dis ptratio chas indus tax 1stat
## 1 24.0 65.2 4.0900
                         15.3
                                  0 2.31 296 4.98
## 2 21.6 78.9 4.9671
                         17.8
                                 0 7.07 242 9.14
                                 0 7.07 242 4.03
## 3 34.7 61.1 4.9671
                         17.8
## 4 33.4 45.8 6.0622
                         18.7
                                 0 2.18 222 2.94
## 5 36.2 54.2 6.0622
                         18.7
                                 0 2.18 222 5.33
## 6 28.7 58.7 6.0622
                         18.7
                                 0 2.18 222 5.21
library(GGally)
ggpairs(Boston.reg) + theme(axis.text.x = element_text(angle = 45, hjust = 1))
```



Describe the overall association between variables.

Stepwise selection (approach, weaknesses, alternatives)

Full model

```
fit.full <- lm(medv ~ age + dis + ptratio + chas + indus + tax + lstat, data = Boston)
msummary(fit.full)
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 58.4021721 2.6119146 22.360 < 2e-16 ***
## age
              0.0020650 0.0143719 0.144 0.885807
## dis
              -1.0708948  0.1966344  -5.446  8.10e-08 ***
## ptratio
             ## chas
              3.4915589 0.9878544 3.534 0.000447 ***
## indus
             -0.2320387  0.0643231  -3.607  0.000341 ***
             -0.0002464 0.0022170 -0.111 0.911539
## tax
             -0.8524239 0.0475568 -17.924 < 2e-16 ***
## 1stat
##
## Residual standard error: 5.491 on 498 degrees of freedom
## Multiple R-squared: 0.6485, Adjusted R-squared: 0.6436
## F-statistic: 131.3 on 7 and 498 DF, p-value: < 2.2e-16
fit.red1 <- lm(medv ~ age + dis + ptratio + chas + indus + lstat, data = Boston)
msummary(fit.red1)
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 58.41054
                         2.60825 22.395 < 2e-16 ***
              0.00206
## age
                         0.01436 0.143 0.885954
## dis
              -1.06990 0.19624 -5.452 7.84e-08 ***
              -1.01789
## ptratio
                         0.12599 -8.079 4.95e-15 ***
                         0.98460 3.554 0.000416 ***
## chas
              3.49901
## indus
              -0.23533
                         0.05706 -4.124 4.36e-05 ***
             -0.85309
                         0.04713 -18.102 < 2e-16 ***
## 1stat
##
## Residual standard error: 5.485 on 499 degrees of freedom
## Multiple R-squared: 0.6485, Adjusted R-squared: 0.6443
## F-statistic: 153.5 on 6 and 499 DF, p-value: < 2.2e-16
```

```
fit.red2 <- lm(medv ~ dis + ptratio + chas + indus + lstat, data = Boston)
msummary(fit.red2)</pre>
```

```
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 58.56949
                          2.35903 24.828 < 2e-16 ***
## dis
              -1.08478
                         0.16642 -6.518 1.74e-10 ***
              -1.01785
## ptratio
                         0.12587 -8.087 4.66e-15 ***
## chas
              3.50877
                         0.98129 3.576 0.000383 ***
## indus
              -0.23458
                         0.05677 -4.132 4.21e-05 ***
## lstat
              -0.85080
                         0.04430 -19.205 < 2e-16 ***
##
## Residual standard error: 5.48 on 500 degrees of freedom
## Multiple R-squared: 0.6485, Adjusted R-squared: 0.645
## F-statistic: 184.5 on 5 and 500 DF, p-value: < 2.2e-16
```

Final model:

Coefficient for chas

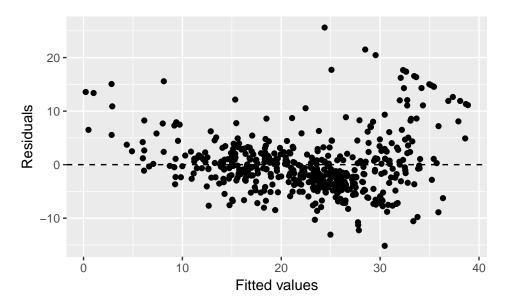
Coefficient for dis

Model validation

- 1. Linearity
- 2. Nearly normal residuals
- 3. Constant variability

Linearity

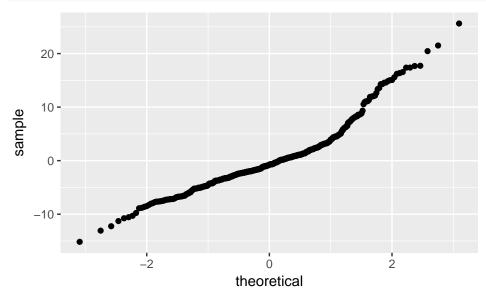
```
qplot(x = .fitted, y = .resid, data = fit.red2) +
  geom_hline(yintercept = 0, linetype = "dashed") +
  xlab("Fitted values") +
  ylab("Residuals")
```



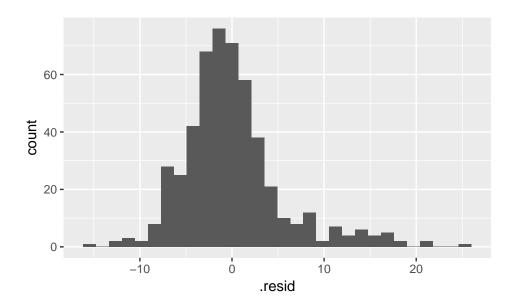
How else to assess linearity?

Nearly normal residuals





qplot(x = .resid, data = fit.red2, geom = "histogram")



Constant variability

```
qplot(x = .fitted, y = .resid, data = fit.red2) +
  geom_hline(yintercept = 0, linetype = "dashed") +
  xlab("Fitted values") +
  ylab("Residuals")
```

