IS5 in R: Linear Regression (Chapter 7)

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Introduction and background

This document is intended to help describe how to undertake analyses introduced as examples in the Fifth Edition of *Intro Stats* (2018) by De Veaux, Velleman, and Bock. More information about the book can be found at http://wps.aw.com/aw_deveaux_stats_series. This file as well as the associated R Markdown reproducible analysis source file used to create it can be found at http://nhorton.people.amherst.edu/is5.

This work leverages initiatives undertaken by Project MOSAIC (http://www.mosaic-web.org), an NSF-funded effort to improve the teaching of statistics, calculus, science and computing in the undergraduate curriculum. In particular, we utilize the mosaic package, which was written to simplify the use of R for introductory statistics courses. A short summary of the R needed to teach introductory statistics can be found in the mosaic package vignettes (http://cran.r-project.org/web/packages/mosaic). A paper describing the mosaic approach was published in the R Journal: https://journal.r-project.org/archive/2017/RJ-2017-024.

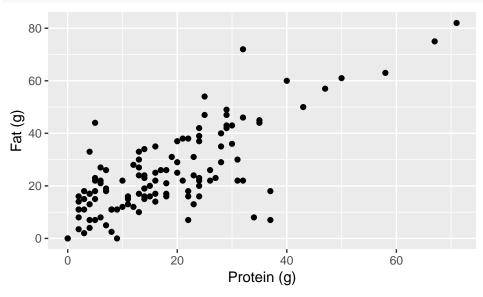
Chapter 7: Linear Regression

```
library(mosaic)
library(readr)
library(janitor)
# Figure 7.1
BurgerKing <- read_csv("http://nhorton.people.amherst.edu/is5/data/Burger_King_items.csv") %>%
  clean names()
## Parsed with column specification:
## cols(
##
     Item = col_character(),
##
     Serving.size = col_integer(),
##
     Calories = col_integer(),
     Fat.Cal = col_integer(),
##
##
     `Protein(g)` = col integer(),
##
     `Fat(g)` = col_double(),
##
     `Sat.Fat(g)` = col_double(),
     `Trans.fat(g)` = col_double(),
##
##
     `Chol(mg)` = col integer(),
     `Sodium(mg)` = col_integer(),
##
##
     `Carbs(g)` = col_integer(),
##
     `Fiber(g)` = col_integer(),
     `Sugar(g)` = col_integer(),
##
     Meat = col_integer(),
##
##
     Breakfast = col_integer(),
##
     `Not Breakfast` = col_integer(),
##
     CarbsxMeat = col_integer()
## )
```

By default, read_csv() prints the variable names. These messages can be suppressed using the message=FALSE code chunk option to save space and improve readability.

Here we use the clean_names() function from the janitor package to sanitize the names of the columns (which would otherwise contain special characters or whitespace).

```
gf_point(fat_g ~ protein_g, data = BurgerKing) %>%
gf_labs(x = "Protein (g)", y = "Fat (g)")
```



Section 7.1: Least Squares: The Line of "Best Fit"

See display on page 197.

36.70931

We can calculate the residual for a particular value with 31 grams of protein by creating an function called burgerfun using the makeFun() function.

```
burgerlm <- lm(fat_g ~ protein_g, data = BurgerKing)
burgerfun <- makeFun(burgerlm)
burgerfun(protein_g = 31)
## 1</pre>
```

Section 7.2: The Linear Model

```
coef(burgerlm)

## (Intercept) protein_g
## 8.4021494 0.9131343

burgerfun(protein_g = 0)

## 1
## 8.402149

burgerfun(32) - burgerfun(31)

## 1
## 0.9131343
```

msummary(burgerlm) Estimate Std. Error t value Pr(>|t|) ## ## (Intercept) 8.40215 1.60400 5.238 7.02e-07 *** ## protein_g 0.91313 0.07177 12.723 < 2e-16 *** ## ## Residual standard error: 10.57 on 120 degrees of freedom ## Multiple R-squared: 0.5743, Adjusted R-squared: 0.5707 ## F-statistic: 161.9 on 1 and 120 DF, p-value: < 2.2e-16 Example 7.1: A Linear Model for Hurricanes Hurricanes <- read_csv("http://nhorton.people.amherst.edu/is5/data/Hurricanes_2015.csv") %>% clean_names() ## Parsed with column specification: ## Name = col_character(), ## Year = col_integer(), `Max.Wind.Speed(kts)` = col_integer(), ## `Central.Pressure(mb)` = col_integer(), ## Category = col_integer() ## ##) gf_point(max_wind_speed_kts ~ central_pressure_mb, data = Hurricanes) %>% gf_labs(x = "Central Pressure (mb)", y = "Max Wind Speed (kts)") ## Warning: Removed 7 rows containing non-finite values (stat_lm). ## Warning: Removed 7 rows containing missing values (geom_point). 150 -Max Wind Speed (kts) 125 -100 -75 **-**50 -920 940 960 980 1000

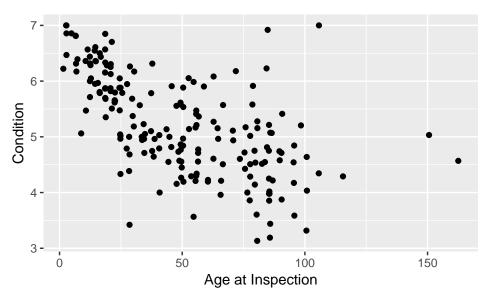
The function generates a warning because some of the data are missing: this output can be suppressed by adding warning=FALSE as an option in this code chunk.

Central Pressure (mb)

Section 7.3: Finding the Least Squares Line

Example 7.2: Finding the Regression Equation

```
favstats(~ protein_g, data = BurgerKing)
## min Q1 median
                     Q3 max
                                            sd
                                                 n missing
      0 7
           15.5 24.75 71 17.93443 13.38911 122
favstats(~ fat_g, data = BurgerKing)
## min
           Q1 median Q3 max
                                mean
                                           sd
                                                 n missing
##
     0 14.25
                  22 33 82 24.77869 16.13362 122
sx <- sd(~ protein_g, data = BurgerKing)</pre>
SX
## [1] 13.38911
sy <- sd(~ fat_g, data = BurgerKing)</pre>
## [1] 16.13362
r <- cor(protein_g ~ fat_g, data = BurgerKing)
r # same as cor(fat_g ~ protein_g)!
## [1] 0.7578003
r*sy/sx
## [1] 0.9131343
coef(burgerlm)[2]
## protein_g
## 0.9131343
Step-by-Step Example: Calculating a Regression Equation
TompkinsBridges <-
 read_csv("http://nhorton.people.amherst.edu/is5/data/Tompkins_county_bridges_2016.csv") %>%
 clean_names()
## Parsed with column specification:
## cols(
##
    Municipality = col_character(),
    Location = col_character(),
##
##
    Route = col_character(),
##
     Owner = col_character(),
##
    Built = col_integer(),
##
    Date.Inspected = col_character(),
    SD.FO.Status = col_character(),
##
    Condition = col_double(),
##
    YearInspected = col_double(),
##
     AgeAtInspection = col_double()
## )
gf_point(condition ~ age_at_inspection, data = TompkinsBridges) %>%
 gf_labs(x = "Age at Inspection", y = "Condition")
```



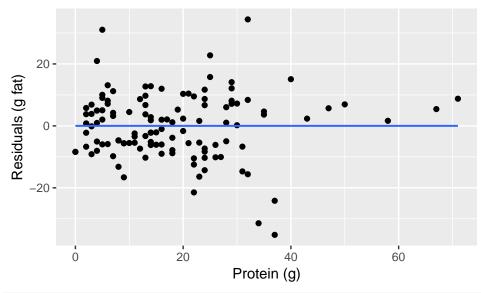
See calculations on page 203.

Section 7.4: Regression to the Mean

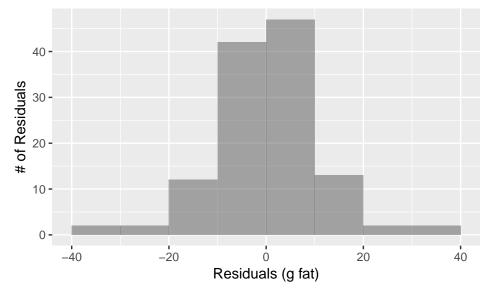
See Figure 7.4 on page 205 to visualize standard deviations.

Section 7.5: Examining the Residuals

```
msummary(burgerlm)
               Estimate Std. Error t value Pr(>|t|)
                                     5.238 7.02e-07 ***
## (Intercept)
               8.40215
                           1.60400
## protein_g
                0.91313
                           0.07177 12.723 < 2e-16 ***
##
## Residual standard error: 10.57 on 120 degrees of freedom
## Multiple R-squared: 0.5743, Adjusted R-squared: 0.5707
## F-statistic: 161.9 on 1 and 120 DF, \, p-value: < 2.2e-16
#Figure 7.5 , page 207
gf_point(resid(burgerlm) ~ protein_g, data = BurgerKing) %>%
  gf_lm() %>%
  gf_labs(x = "Protein (g)", y = "Residuals (g fat)")
```



```
# Figure 7.6
gf_histogram(~ resid(burgerlm), binwidth = 10, center = 5) %>%
gf_labs(x = "Residuals (g fat)", y = "# of Residuals")
```



Section 7.6: R^2 -The Proportion of Variation Accounted for by the Model

```
rsquared(burgerlm)
```

[1] 0.5742613

Section 7.7: Regression Assumptions and Conditions