Intro stats with mosaic

agformula version

Loading packages

library (mosaic)

Essential R syntax

Names in R are case sensitive Function and arguments

rflip(10)

Optional arguments

rflip(10, prob = 0.8)

Assignment

 $x \leftarrow rflip(10, prob = 0.8)$

Getting help on any function

help(mean)

Arithmetic operations

basic operations exponentiation grouping () square root sqrt(x) absolute value abs(x)logarithm, base 10 log10(x)log(x)natural logarithm, base e exponential function e^x exp(x)

Logical operators

is equal to (note double equal sign)

factorial (k) $k! = k(k-1) \dots 1$

- is not equal to
- is less than
- is less than or equal to
- is greater than
- is greater than or equal to
- A & B ("A and B") is TRUE if both A and B are TRUE
- A | B ("A or B") is TRUE if one or both of A and B are TRUE

%in% inclusion; for example

"C" %in% c("A", "B") is FALSE

Formula interface

Use for graphics, statistics, inference, and modeling operations.

 $goal(y \sim x, data = mydata)$ Read as "Calculate goal for v using mydata "broken down by" x, or "modeled by" x.

mean(age ~ sex, data = HELPrct)

For graphics:

 $goal(y \sim x \mid z, data = mydata,$ $color = \sim w)$

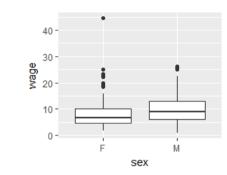
y: y-axis variable (optional)

x: x-axis variable (required)

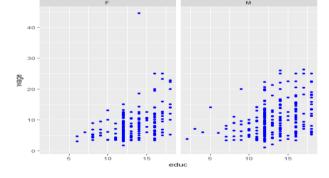
z: panel-by variable (optional)

w: color-by formula (optional)

gf boxplot(wage ~ sex, data = CPS85)



gf point(wage ~ educ | sex, data = CPS85, color = "blue")



gf point(wage ~ educ, data = CPS85, color = ~ sex)



Examining data

Print short summary of all variables inspect(HELPrct) Number of rows and columns

dim(HELPrct) nrow (HELPrct) ncol (HELPrct)

Print first rows or last rows

head(KidsFeet) tail(KidsFeet, 10)

Names of variables

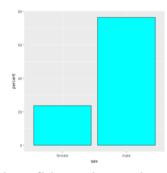
names (HELPrct)

One categorical variable

Counts by category

tally(~ sex, data = HELPrct) Percentages by category tally(~ sex, data = HELPrct, format = "percent")

Bar graph of percentages gf percents(~ sex, data = HELPrct, fill = "cyan", color = "black")



Tests and confidence intervals

Exact test

result1 <-

binom.test(~ homeless == "homeless", data = HELPrct)

Approximate test (large samples)

result2 <-

prop.test(~ homeless == "homeless", data = HELPrct, alternative = "less", p = 0.4)

Extract confidence intervals and *p*-values confint(result1) pval(result2)

One quantitative variable

Make output more readable options(digits = 3)

Compute summary statistics

mean(~ cesd, data = HELPrct)

Other summary statistics work similarly

median() iqr() max() min() fivenum() sd() var() sum()

Table of summary statistics

favstats(~ cesd, data = HELPrct)

Summary statistics by group

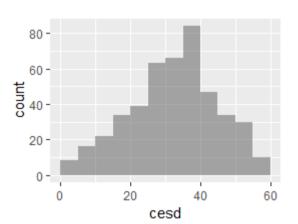
favstats(cesd ~ sex, data = HELPrct)

Quantiles

quantile (~ cesd, data = HELPrct, prob = c(0.25, 0.5, 0.8))

Histogram

gf histogram(~ cesd, data = HELPrct, binwidth = 5, center = 2.5)



Normal probability plot gf qq(~ cesd, data = HELPrct)

Density plot

gf dens(~ cesd, data = HELPrct, color = "blue", size = 1.25)

One-sample *t*-test result <- t test(~ cesd, data = HELPrct, mu = 34)

Extract confidence intervals and *p*-values confint(result) pval(result)

Paired *t*-test

t test(extra ~ group, data = sleep, paired = TRUE)

Data wrangling

Drop, rename, or reorder variables df <- select(HELPrct,</pre> c(id, age, gender = sex)) Create new variables from existing ones KidsFeet <- mutate(KidsFeet,</pre> width in = 0.394 * width) Extract specific rows from data girls feet <- filter(KidsFeet,</pre> sex == "G")Sort data rows by value in column df <- arrange(KidsFeet, length)</pre> Compute summary statistics by group group by(KidsFeet, sex) %>% summarize(mean width = mean(width)) For more, see Tidyverse cheatsheet

Importing data

Import data from file or URL

MustangPrice < read.file("C:/MustangPrice.csv")

Note: R uses forward slashes in file paths

kidsfeet < read.file("http://www.mosaic web.org/go/datasets/kidsfeet.csv")</pre>

Randomization and simulation

Fix random number sequence set.seed(42) Toss coins rflip(10) # default prob is 0.5 Do something repeatedly do(5) * rflip(10, prob = 0.75)Draw a simple random sample sample(LETTERS, 10) deal(Cards, 5) # poker hand Resample with replacement Small <- sample(KidsFeet, 10)</pre> resample(Small) Random permutation (shuffling) shuffle (Cards) Random values from distributions rbinom(5, size = 10, prob = 0.7)rnorm(5, mean = 10, sd = 2)

Two categorical variables

Contingency table with margins

tally(~ substance + sex,
 data = HELPrct, margins = TRUE)

Percentages by column

tally(~ sex | substance,
 data = HELPrct,
 format = "percent")

Mosaic plot

my_tbl <- tally(substance ~ sex,
 data = HELPrct)

mosaicplot(my_tbl, color = TRUE)

substance

Test for proportions (approximate)
prop.test(homeless ~ sex,
 success = "homeless",
 data = HELPrct)

Distributions

Normal distribution function pnorm(13, mean = 10, sd = 2)Normal distribution function with graph xpnorm(1.645, mean = 0, sd = 1)Normal distribution quantiles qnorm(0.95) # mean = 0, sd = 1Normal distribution quantiles with graph xqnorm(0.85, mean = 10, sd = 2)Binomial density function ("size" means *n*) dbinom(5, size = 8, prob = 0.65)Binomial distribution function pbinom(5, size = 8, prob = 0.65)Central portion of distribution cdist("norm", 0.95) cdist("t", c(0.90, 0.99), df = 5)Plotting distributions plotDist("binom", size = 8, prob = 0.65, xlim = c(-1, 9)plotDist("norm", mean = 10, sd = 2

Two quantitative variables

data = HELPrct)
msummary(cesdmodel)

Prediction

lm_fun <- makeFun(cesdmodel)

lm_fun(mcs = 35)

Extract useful quantities
anova(cesdmodel)
coef(cesdmodel)
confint(cesdmodel)
rsquared(cesdmodel)

Diagnostics; plot residuals

gf_dhistogram(~resid(cesdmodel))

gf_qq(~resid(cesdmodel))</pre>
Diagnostics; plot residuals vs. fitted

Simple linear regression

cesdmodel <- lm(cesd ~ mcs,

Categorical response, quantitative predictor

gf point(resid(cesdmodel) ~

gf lm(size = 2)

fitted(cesdmodel)) %>%

Logistic regression

logit_mod <- glm(homeless ~ age,
 data = HELPrct,
 family = binomial)

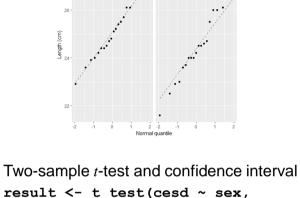
msummary(logit_mod)

Odds ratios and confidence intervals

exp(coef(logit_mod))

exp(confint(logit_mod))

Quantitative response, categorical predictor



data = HELPrct)

```
result # view results
confint(result)
pval(result)

More than two levels (Analysis of variance)
Numeric and graphic summaries
favstats(cesd ~ substance,
    data = HELPrct)

gf_boxplot(cesd ~ substance,
    data = HELPrct)

Fit and summarize model
mod <- lm(age ~ substance,
    data = HELPrct)
anova(mod)

Which differences are significant?
mplot(TukeyHSD(mod))</pre>
```

