

Intro stats with mosaic

ggformula 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 <- rflip(10, prob = 0.8)
```

Getting help on any function

```
help(mean)
```

Arithmetic operations

<code>+</code>	<code>-</code>	<code>*</code>	<code>/</code>	basic operations
<code>^</code>				exponentiation
<code>()</code>				grouping
<code>sqrt(x)</code>				square root
<code>abs(x)</code>				absolute value
<code>log10(x)</code>				logarithm, base 10
<code>log(x)</code>				natural logarithm, base e
<code>exp(x)</code>				exponential function e^x
<code>factorial(k)</code>				$k! = k(k - 1) \dots 1$

Logical operators

<code>==</code>	is equal to (note double equal sign)
<code>!=</code>	is not equal to
<code><</code>	is less than
<code><=</code>	is less than or equal to
<code>></code>	is greater than
<code>>=</code>	is greater than or equal to
<code>&</code>	<code>A & B</code> ("A and B") is TRUE if both A and B are TRUE
<code> </code>	<code>A B</code> ("A or B") is TRUE if one or both of A and B are TRUE
<code>%in%</code>	inclusion; for example <code>"C" %in% c("A", "B")</code> is FALSE

Formula interface

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

```
goal(y ~ x, data = mydata)
Read as "Calculate goal for y using
mydata "broken down by" x, or
"modeled by" x.
mean(age ~ sex, data = HELPrc)
```

For graphics:

```
goal(y ~ x | z, data = mydata,
      color = ~ w)
```

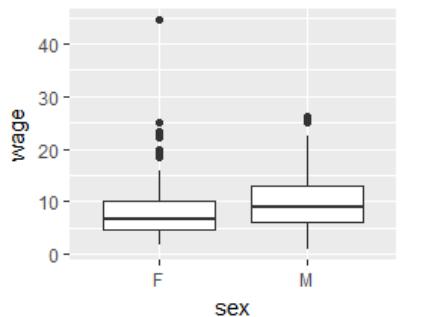
`y`: y-axis variable (*optional*)

`x`: x-axis variable (*required*)

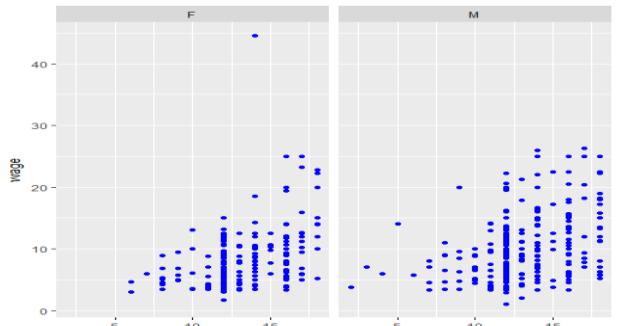
`z`: panel-by variable (*optional*)

`w`: color-by formula (*optional*)

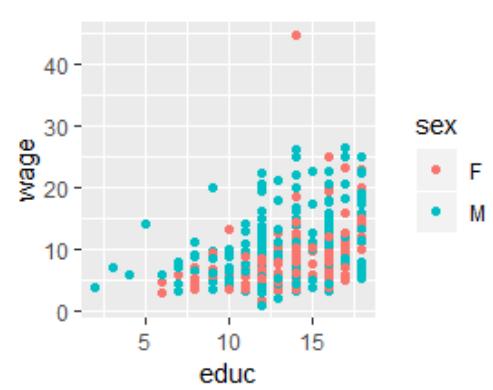
```
gf_boxplot(wage ~ sex,
           data = CPS85)
```



```
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(HELPrc)
```

Number of rows and columns

```
dim(HELPrc)
```

```
nrow(HELPrc)
```

```
ncol(HELPrc)
```

Print first rows or last rows

```
head(KidsFeet)
```

```
tail(KidsFeet, 10)
```

Names of variables

```
names(HELPrc)
```

One categorical variable

Counts by category

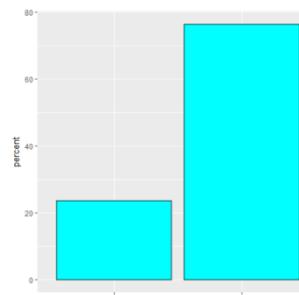
```
tally(~ sex, data = HELPrc)
```

Percentages by category

```
tally(~ sex, data = HELPrc,
      format = "percent")
```

Bar graph of percentages

```
gf_percents(~ sex,
            data = HELPrc, fill = "cyan",
            color = "black")
```



Tests and confidence intervals

Exact test

```
result1 <-
binom.test(~ homeless ==
"homeless", data = HELPrc)
```

Approximate test (large samples)

```
result2 <-
prop.test(~ homeless ==
"homeless", data = HELPrc,
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 = HELPrc)
```

Other summary statistics work similarly

```
median() iqr() max() min()
```

```
fivenum() sd() var() sum()
```

Table of summary statistics

```
favstats(~ cesd, data = HELPrc)
```

Summary statistics by group

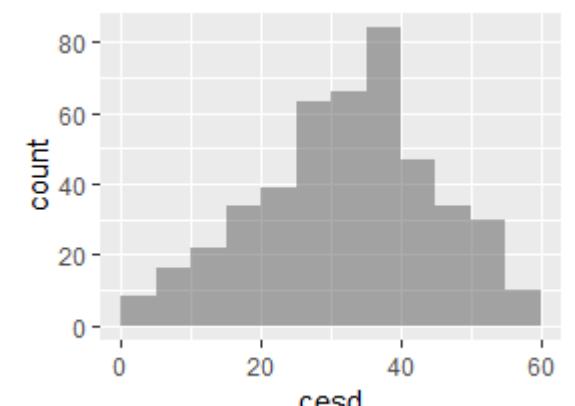
```
favstats(cesd ~ sex,
         data = HELPrc)
```

Quantiles

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

Histogram

```
gf_histogram(~ cesd,
            data = HELPrc, binwidth = 5,
            center = 2.5)
```



Normal probability plot

```
gf_qq(~ cesd, data = HELPrc)
```

Density plot

```
gf_dens(~ cesd, data = HELPrc,
        color = "blue", size = 1.25)
```

One-sample t-test

```
result <- t_test(~ cesd,
                 data = HELPrc, 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(HELPrc, c(id, age, gender = sex))  
  
Create new variables from existing ones  
KidsFeet <- mutate(KidsFeet, width_in = 0.394 * width)  
  
Extract specific rows from data  
girls_feet <- filter(KidsFeet, sex == "G")  
  
Sort data rows by value in column  
df <- arrange(KidsFeet, length)  
  
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")
```

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)  
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 = HELPrc, margins = TRUE)  
  
Percentages by column  
tally(~ sex | substance,  
      data = HELPrc,  
      format = "percent")  
  
Mosaic plot  
my_tbl <- tally(substance ~ sex,  
                 data = HELPrc)  
mosaicplot(my_tbl, color = TRUE)  
  

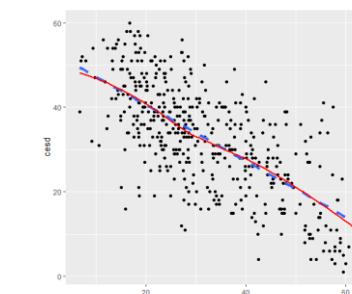
```

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

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 = 1  
  
Normal 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

```
Correlation coefficient  
cor(cesd ~ mcs, data = HELPrc)  
  
Scatterplot with regression line and smooth  
gf_point(cesd ~ mcs,  
          data = HELPrc) %>%  
  gf_lm(size = 1.5, linetype =  
    "dashed") %>%  
  gf_smooth(color = "red")  
  

```

```
Simple linear regression  
cesdmodel <- lm(cesd ~ mcs,  
                  data = HELPrc)  
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))  
  
Diagnostics; plot residuals vs. fitted  
gf_point(resid(cesdmodel) ~  
          fitted(cesdmodel))
```

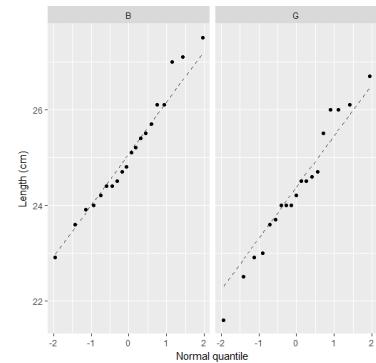
Categorical response, quantitative predictor

```
Logistic regression  
logit_mod <- glm(homeless ~ age,  
                   data = HELPrc,  
                   family = binomial)  
msummary(logit_mod)  
  
Odds ratios and confidence intervals  
library(broom)  
tidy(logit_mod, conf.int = TRUE,  
     exponentiate = TRUE)
```

Quantitative response, categorical predictor

Two-level predictor: two-sample *t* test

```
Numeric summaries  
favstats(~length | sex,  
         data = KidsFeet)  
  
Graphic summaries  
gf_qq(~ length | sex,  
      data = KidsFeet) %>%  
  gf_qqline() %>%  
  gf_labs(x = "Normal quantile",  
          y = "Length (cm)")
```



Two-sample *t*-test and confidence interval

```
result <- t_test(cesd ~ sex,  
                  data = HELPrc)  
result # view results  
confint(result)  
pval(result)
```

More than two levels (Analysis of variance)

```
Numeric and graphic summaries  
favstats(cesd ~ substance,  
         data = HELPrc)  
gf_boxplot(cesd ~ substance,  
           data = HELPrc)
```

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

Which differences are significant?

```
mplot(TukeyHSD(mod))
```

