

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

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

Logical operators

| | |
|------|---|
| == | is equal to (note double equal sign) |
| != | is not equal to |
| < | is less than |
| <= | is less than or equal to |
| > | is greater than |
| >= | is greater than or equal to |
| & | A & B is TRUE if both A and B are TRUE |
| | A 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 ~ x, data = mydata)
```

Read as "Calculate goal for y using mydata "broken down by" x, or "modeled by" x.

```
mean(age ~ sex, data = HELPrct)
```

For graphics:

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

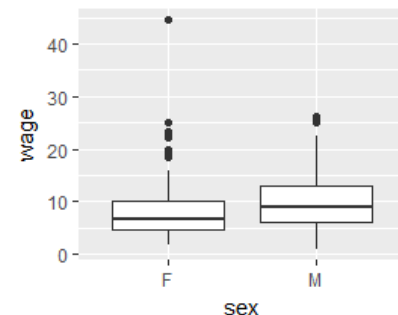
y : y-axis variable (*optional*)

x : x-axis variable (*required*)

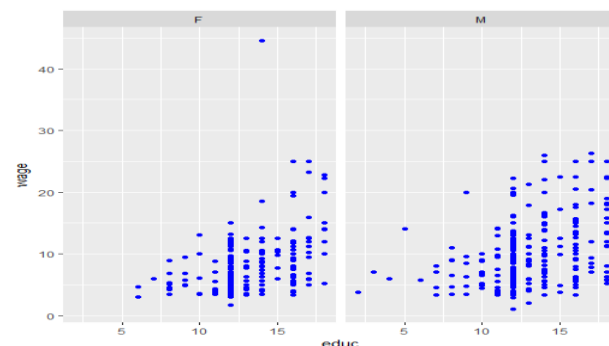
z : panel-by variable (*optional*)

w : color-by formula (*optional*)

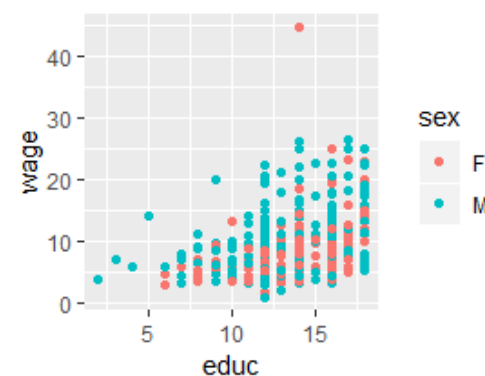
```
gf_boxplot(wage ~ sex, data = CPS85)
```



```
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(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

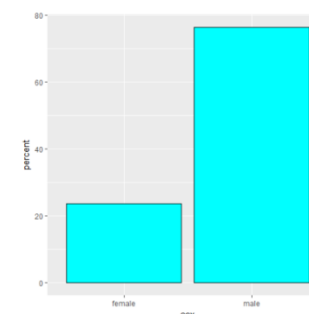
```
tally(~ sex, data = HELPrct)
```

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

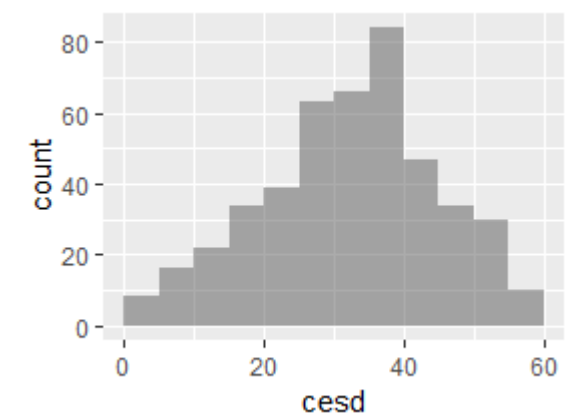
```
favstats(cesd ~ sex, data = HELPrct)
```

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,
  c(id, age, sex))

Create new variables from existing ones
KidsFeet <- mutate(KidsFeet,
  width_in = 0.394 * width)

Retain 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!
Dome <-
  read.file("http://www.mosaic-
  web.org/go/datasets/Dome.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 = HELPrct, margins = TRUE)

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

Mosaic plot
my_tbl <- tally(sex ~ substance,
  data = HELPrct)
mosaicplot(my_tbl, color = TRUE)
```



```
Chi-square test
xchisq.test(~ substance + sex,
  data = HELPrct,
  correct = FALSE)
```

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)

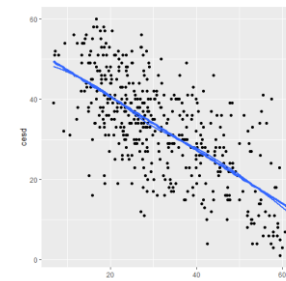
Central portion of distribution
cdist("norm", 0.95)
cdist("t", c(0.90, 0.99), df = 5)

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 = HELPrct)

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



```
Simple linear regression
cesdmodel <- lm(cesd ~ mcs,
  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))

Diagnostics; plot residuals vs. fitted
gf_point(resid(cesdmodel) ~
  fitted(cesdmodel)) %>%
  gf_lm(size = 2)
```

Categorical response, quantitative predictor

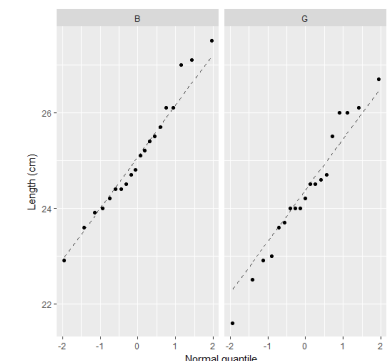
```
Logistic regression
logit_mod <- glm(homeless ~ age,
  data = HELPrct,
  family = binomial)
msummary(logit_mod)

Odds ratios and confidence intervals
exp(coef(logit_mod))
exp(confint(logit_mod))
```

Quantitative response, categorical predictor

```
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)")
```



```
gf_boxplot(cesd ~ substance,
  data = HELPrct)
```

```
Two-sample t-test and confidence interval
result <- t_test(cesd ~ sex,
  data = HELPrct)
result # view results
confint(result)
pval(result)
```

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

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

Which differences are significant?
mplot(TukeyHSD(mod))
```

