

Intro stats with mosaic

(lattice version)

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)`

Loading packages

`library(mosaic)`

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> is TRUE if both <code>A</code> and <code>B</code> are TRUE
<code> </code>	<code>A B</code> is TRUE if one or both of <code>A</code> and <code>B</code> are TRUE
<code>%in%</code>	includes; 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, groups = w,
      data = mydata)
```

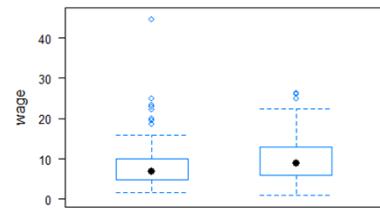
`y` : y-axis variable (*optional*)

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

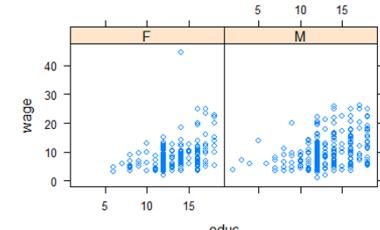
`z` : panel-by variable (*optional*)

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

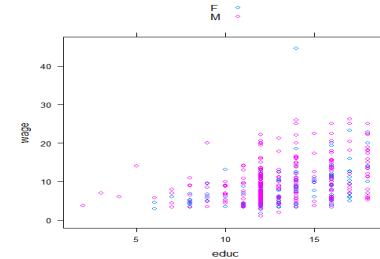
```
bwplot(wage ~ sex, data = CPS85)
```



```
xyplot(wage ~ educ | sex,
       data = CPS85)
```



```
xyplot(wage ~ educ,
       groups = sex, data = CPS85,
       auto.key = TRUE)
```



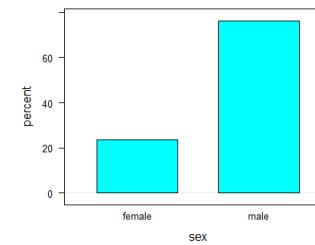
One categorical variable

Counts by category

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

Percentages by category

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



Tests and confidence intervals

Exact test

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

Approximate test (large samples)

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

Extract confidence intervals and p-values

```
confint(result1)
pval(result2)
```

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 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

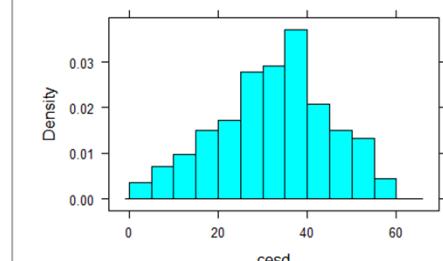
```
favstats(cesd ~ sex,
  data = HELPrc)
```

Quantiles

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

Histogram

```
histogram(~ cesd, width = 5,
  center = 2.5, data = HELPrc)
```



Normal probability plot

```
qqmath(~ cesd, dist = "qnorm",
  data = HELPrc)
```

Density plot

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

Dot plot

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

One-sample t-test

```
result <- t.test(~ cesd, mu =
  34, data = HELPrc)
```

Extract confidence intervals and p-values

```
confint(result)
```

```
pval(result)
```

Two categorical variables

Contingency table with margins

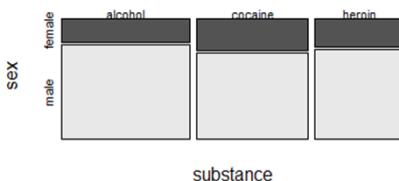
```
tally(~ substance + sex,  
      margins = TRUE,  
      data = HELPrc)
```

Percentages by column

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

Mosaic plot

```
mosaicplot(~ substance + sex,  
           color = TRUE, data = HELPrc)
```



Chi-square test

```
xchisq.test(~ substance + sex,  
            data = HELPrc,  
            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)
```

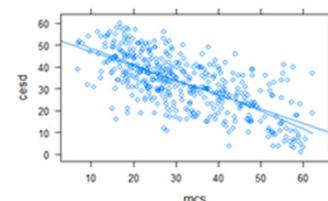
Two quantitative variables

Correlation coefficient

```
cor(cesd ~ mcs, data = HELPrc)
```

Scatterplot with regression line and smooth

```
xyplot(cesd ~ mcs,  
       type = c("p", "r", "smooth"),  
       data = HELPrc)
```



Simple linear regression

```
cesdmodel <- lm(cesd ~ mcs,  
                  data = HELPrc)  
msummary(cesdmodel)
```

Prediction

```
lmfunction <- makeFun(cesdmodel)  
lmfunction(mcs = 35)
```

Extract useful quantities

```
anova(cesdmodel)  
coef(cesdmodel)  
confint(cesdmodel)  
rsquared(cesdmodel)
```

Diagnostics; plot residuals

```
histogram(~resid(cesdmodel),  
          density = TRUE)  
qqmath(~resid(cesdmodel))
```

Diagnostics; plot residuals vs. fitted

```
xyplot(resid(cesdmodel) ~  
       fitted(cesdmodel),  
       type = c("p", "smooth", "r"))
```

Categorical response, quantitative predictor

Logistic regression

```
logit_mod <-  
  glm(homeless ~ age + female,  
       family = binomial, data = HELPrc)  
msummary(logitmod)
```

Odds ratios and confidence intervals

```
exp(coef(logit_mod))  
exp(confint(logit_mod))
```

Data management

From dplyr package

Drop or reorder variables

```
select()
```

Create new variables from existing ones

```
mutate()
```

Retain specific rows from data

```
filter()
```

Sort data rows

```
arrange()
```

Compute summary statistics by group

```
group_by()
```

```
summarize()
```

Merge data tables

```
left_join()
```

```
inner_join()
```

Importing data

Import file from computer 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)
```

Tossing 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)
```

Quantitative response, categorical predictor

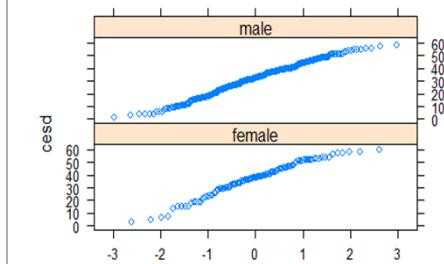
Two-level predictor: two-sample t test

Numeric summaries

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

Comparative normal probability plot

```
qqmath(~cesd | sex, data = HELPrc,  
      layout = c(1, 2)) # also bwplot
```



Dotplot for smaller samples

```
xyplot(sex ~ length, alpha = 0.6,  
      cex = 1.4, data = KidsFeet)
```

Two-sample t -test and confidence interval

```
result <- t.test(cesd ~ sex,  
                  var.equal = FALSE, data = HELPrc)  
confint(result)
```

More than two levels: Analysis of variance

Numeric summaries

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

Graphic summaries

```
bwplot(cesd ~ substance, pch = "|",  
       data = HELPrc)
```

Fitt and summarize model

```
modsubstance <- lm(cesd ~ substance,  
                     data = HELPrc)  
anova(modsubstance)
```

Which differences are significant?

```
pairwise <- TukeyHSD(modsubstance)  
mplot(pairwise)
```

95% family-wise confidence level

