

# Intro stats with mosaic

ggformula 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

+	-	*	/	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
&	<b>A &amp; B</b> is <b>TRUE</b> if both <b>A</b> and <b>B</b> are <b>TRUE</b>
	<b>A   B</b> is <b>TRUE</b> if one or both of <b>A</b> and <b>B</b> are <b>TRUE</b>
%in%	includes; for example "C" %in% c("A", "B") is <b>FALSE</b>

## 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, color = ~ w,
      data = mydata)
```

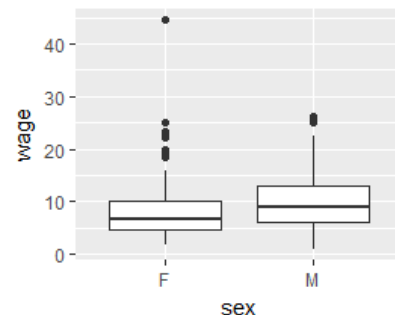
**y** : y-axis variable (*optional*)

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

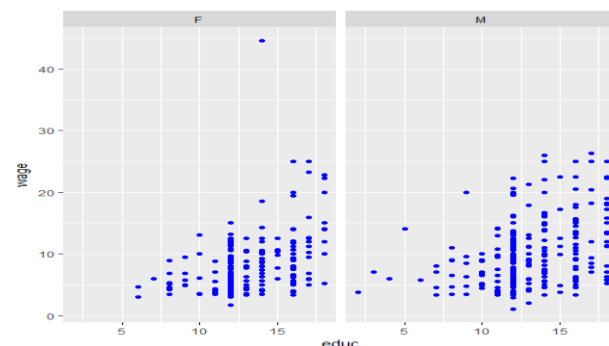
**z** : panel-by variable (*optional*)

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

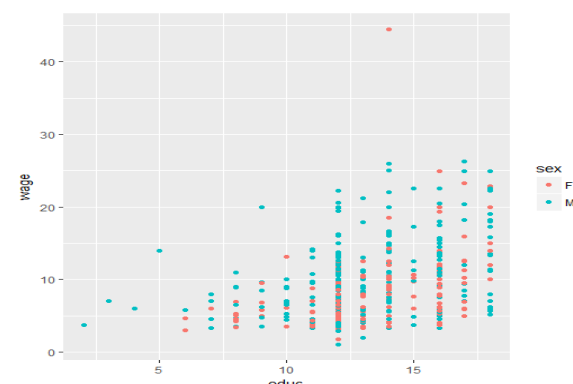
```
gf_boxplot(wage ~ sex,
            data = CPS85)
```



```
gf_point(wage ~ educ | sex,
          data = CPS85, color = "blue")
```



```
gf_point(wage ~ educ,
          color = ~ sex, data = CPS85)
```



## 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)
```

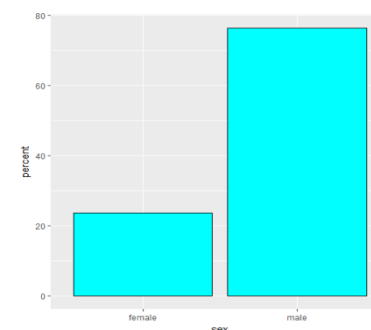
## One categorical variable

Counts by category

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

Percentages by category

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

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

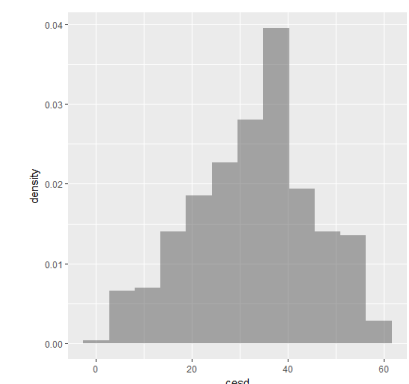
```
favstats(cesd ~ sex,
          data = HELPrct)
```

Quantiles

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

Histogram

```
gf_dhistogram(~ cesd, data =
              HELPrct, bins = 12)
```



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,
                 mu = 34, data = HELPrct)
```

Extract confidence intervals and  $p$ -values

```
confint(result)
pval(result)
```

## Data manipulation

From `dplyr` package  
For details, see [Tidyverse cheatsheet](#)

Drop, rename, 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()`

## 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)`

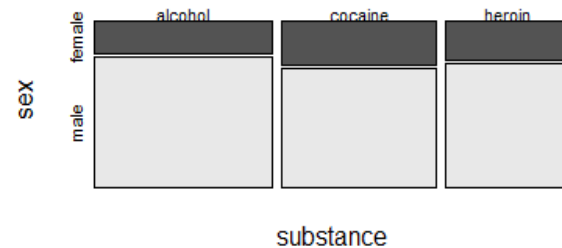
Random values from distributions  
`rbinom(5, size = 10, prob = 0.7)`  
`rnorm(5, mean = 10, sd = 2)`

## Two categorical variables

Contingency table with margins  
`tally(~ substance + sex,  
 margins = TRUE,  
 data = HELPrct)`

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

Mosaic plot  
`mosaicplot(~ substance + sex,  
 color = TRUE, data = HELPrct)`



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)`

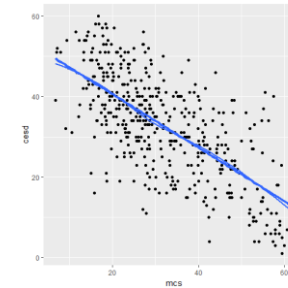
Central portion of distribution  
`cdist("norm", 0.95)`  
`cdist("t", c(0.90, 0.99), df = 5)`

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,  
 family = binomial, data = HELPrct)`  
`msummary(logit_mod)`

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

## Quantitative response, categorical predictor

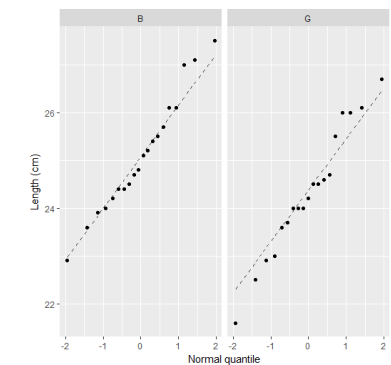
Two-level predictor: two-sample  $t$  test

Numeric summaries

```
favstats(~length | sex,  
  data = KidsFeet)
```

Graphic summaries

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



```
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)
```

More than two levels (Analysis of variance)

Numeric summaries

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

Fit and summarize model

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

Which differences are significant?

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

