## Numerical explorations with R

Waithaka Michael

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### Introduction

- Before analysing any study data, it is common practice to explore the data, to get a broad idea about the phenomenon we are studying.
- Summary statistics are then of interest, such as mean, variability, frequencies.
- It is also of interest to know if missing data are present.
- R provides these measures through the use of the function summary().

- Read in the data
- Datasets in R are typically stored as data frames, which have a matrix structure
- Observations are arranged as rows and variables, either numerical or categorical, are arranged as columns

### Import the dataset

```
data <- read.csv("data/bwmal.csv")</pre>
```

### Get the dimension of the dataset

```
dim(data)
## [1] 791 12
```

### Explore variable names of the dataset

### The dataset at a glance

#### **Explore the structure of the dataset**

```
str(data)
   'data.frame': 791 obs. of 12 variables:
##
   $ X : int 1 2 3 4 5 6 7 8 9 10 ...
   $ matage : int 26 23 18 25 25 21 20 19 32 23 ...
##
##
   $ mheight : num 1.58 1.53 1.54 1.58 1.55 ...
##
   $ gestwks : int 40 40 40 40 40 40 41 38 40 41 ...
##
   $ sex : int 0 0 1 1 1 1 1 1 0 0 ...
##
   $ bweight : num 3.11 2.65 3.41 2.99 3.16 ...
##
   $ smoke : int 0 0 0 0 0 0 0 0 0 ...
   $ pfplacen: int 0 0 0 0 0 0 1 0 0 ...
##
   $ parity : int 3 1 0 2 1 1 0 0 6 0 ...
##
##
   $ workload: int 0 0 0 0 1 1 1 1 0 0 ...
   $ matagegp: int 3 3 1 3 3 2 2 1 4 3 ...
##
##
   $ gestcat : int 2 2 2 2 2 2 2 2 2 2 ...
```

### Viewing data contents of a variable

We can access variables directly by using their names, using the object \$ variable notation

```
data$sex
##
                               0 1 1 0 1 0 1 0 0 1 1
##
##
##
```

### Viewing specific cell contents

To access a certain entry, we most commonly use object[row,column]

```
data[2, 3]
## [1] 1.529
```

### Viewing specific variable contents

all data in variable 5 (sex)

## Viewing specific row/observation contents - all data in row 5

```
data[5, ]

## X matage mheight gestwks sex bweight smoke pfplacen
## 5 5   25  1.555   40  1  3.16  0  0

## parity workload matagegp gestcat
## 5   1   1  3  2
```

## Data in a range - all data in rows 2 and 3, columns 2 and 3

# ....back to the function *summary()*

- This function returns some basic summary statistics, which differ according to the class of the objects that are considered.
- In particular R distinguishes between:
  - numerical vectors: mean,minimum, maximum and quartiles are calculated,
  - factors: frequencies are calculated,
  - character vectors: just the class of the object is returned,
  - ... just try for the rest (but be critical towards the output!).

# The function *summary()*

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```
# summary statistics for continuous variables using the
# function summary()
mydata = data[, c(1:4, 6)]
summary(mydata)
##
        X
                     matage mheight
##
   Min. : 1.0 Min. :13.00 Min. :1.352
## 1st Qu.:198.5 1st Qu.:20.00 1st Qu.:1.506
  Median: 396.0 Median: 23.00 Median: 1.544
##
  Mean :396.0 Mean :23.78
                               Mean :1.543
##
## 3rd Qu.:593.5
                 3rd Qu.:27.00
                               3rd Qu.:1.580
##
   Max. :791.0 Max. :46.00 Max. :1.750
##
  gestwks bweight
##
   Min. :28.00 Min. :0.78
##
   1st Qu.:38.00
                 1st Qu.:2.58
   Median ·39 00
                 Median · 2 90
```

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## Specific functions to summarize the data

- Enables us to see the main characteristics of data before any formal modeling or hypothesis testing
- Particular techniques depends on the type of variable: Continuous or categorical
  - Continuous eg. matage, mheight, gestwks, bweight, parity
  - Categorical eg. smoking status, sex, pfplacen, workload, matagegp, gestcat

### **Examples of data explorations: Continuous variables**

```
min(data$mheight)
## [1] 1.352
max(data$mheight)
## [1] 1.75
```

## Some data explorations: Continuous variables

```
mean(data$mheight)
  [1] 1.543273
var(data$mheight)
  [1] 0.002884892
sd(data$matage)
## [1] 5.139645
median(data$matage)
## [1] 23
```

# More data explorations using function apply()

## Produce the defined summary statistic for continous variables

(mydata.mean = apply(mydata, MARGIN = 2, FUN = mean))

```
## X matage mheight gestwks bweight
## 396.000000 23.782554 1.543273 38.988622 2.900354
(mydata.median = apply(mydata, MARGIN = 2, FUN = median))
##
        X matage mheight gestwks bweight
## 396.000 23.000 1.544 39.000 2.900
(mydata.quantiles = apply(mydata, MARGIN = 2, FUN = sd))
##
           X matage mheight gestwks bweight
## 228.4863234    5.1396448    0.0537112    1.6369536    0.5108436
                                   4日 → 4周 → 4 重 → 4 重 → 9 9 ○
```

### Summarize single categorical variable

```
# freq table for the factor variables
(freq.table.sex = table(mydata2$sex))
##
## 0 1
## 381 410
(freq.table.smoke = table(mydata2$smoke))
##
## 724 67
```

## Cross-tabulation of two categorical variables: 2-Way Frequency Table

```
(mytable <- table(mydata2$sex, mydata2$smoke))</pre>
##
##
## 0 346 35
    1 378 32
##
(mytable <- with(data, table(sex, smoke))) #with command adds
able labels
##
  smoke
## sex 0 1
## 0 346 35
## 1 378 32
```

### Tables of marginal frequencies

```
# sex frequencies (summed over smoke)
margin.table(mytable, 1)
## sex
## 0 1
## 381 410
# smoking status frequencies (summed over sex)
margin.table(mytable, 2)
## smoke
## 0 1
## 724 67
```

### Tables of proportions

```
100 * prop.table(mytable) # cell percentages
## smoke
## sex
## 0 43.742099 4.424779
## 1 47.787611 4.045512
100 * prop.table(mytable, 1) # row percentages
## smoke
## sex
## 0 90.813648 9.186352
##
    1 92.195122 7.804878
```

# column percentages = 100\*prop.table(mytable, 2)

Testing the independence of the row and column variable

```
chisq.test(mytable, correct = FALSE) # chi-square test of in-
depedence
##
##
   Pearson's Chi-squared test
##
## data: mytable
## X-squared = 0.48614, df = 1, p-value = 0.4857
# summary(mytable) - chi-square test of indepedence
# chisq.test(mytable) - chi-square test of indepedence with
# Yates' continuity correction
```

## 3-Way Frequency Table : using xtabs

### Log-linear models for 3-Way Frequency Table

```
library (MASS)
# Mutual Independence: sex, smoking status and matagegp are
# pairwise independent
loglm(~sex + smoke + matagegp, mytable)
## Call:
## loglm(formula = ~sex + smoke + matagegp, data = mytable)
##
## Statistics:
                          X^2 df P(> X^2)
##
## Likelihood Ratio 9.544296 10 0.4813403
                    10.060646 10 0.4351889
## Pearson
```

### Log-linear models for 3-Way Frequency Table

```
# Conditional Independence: sex is independent of smoking
# status, given matagegp.
loglm("sex + smoke + matagegp + sex * matagegp + smoke * matagegp
   mytable)
## Call:
## loglm(formula = ~sex + smoke + matagegp + sex * matagegp +
##
       matagegp, data = mytable)
##
## Statistics:
                         X^2 df P(> X^2)
##
## Likelihood Ratio 0.762029 4 0.9434649
           0.763251 4 0.9433058
## Pearson
```

### Log-linear models for 3-Way Frequency Table

```
# No Three-Way Interaction
loglm("sex + smoke + matagegp + sex * smoke + sex * matagegp +
    smoke * matagegp, mytable)
## Call:
## loglm(formula = ~sex + smoke + matagegp + sex * smoke + sex
##
      matagegp + smoke * matagegp, data = mytable)
##
## Statistics:
                         X^2 df P(> X^2)
##
## Likelihood Ratio 0.2838056 3 0.9630449
          0.2833175 3 0.9631349
## Pearson
```

