Bios 6301: Assignment 6

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Due Thursday, 15 November, 1:00 PM $5^{n=day}$ points taken off for each day late.

30 points total.

Submit a single knitr file (named homework8.rmd), along with a valid PDF output file. Inside the file, clearly indicate which parts of your responses go with which problems (you may use the original homework document as a template). Add your name as author to the file's metadata section. Raw R code/output or word processor files are not acceptable.

Failure to name file homework8.rmd or include author name may result in 5 points taken off.

Question 1

15 points

Install the readxl package and run the following

```
fn <- 'icd10.xlsx'
if(file.access(fn, mode = 4) == -1) {
    url <- "https://www.cdc.gov/nhsn/xls/icd10-pcs-pcm-nhsn-opc.xlsx"
    download.file(url, destfile = fn)
}
dat <- readxl::read_excel(fn, sheet = 2)</pre>
```

1. Show the class of dat. (1 point)

```
class(dat)
```

```
## [1] "tbl df" "tbl" "data.frame"
```

2. Show the methods available for objects of the given class (if there are multiple classes, show methods for all classes). (3 points)

```
methods(class = class(dat)[1])
```

```
[[<-
                                                    [<-
    [1] [
                       [6] $<-
                                                                  initialize
##
                       as.data.frame coerce
                                                    format
                      print
                                                                  slotsFromS3
## [11] Ops
                                     row.names<-
                                                    show
## see '?methods' for accessing help and source code
methods(class = class(dat)[2])
```

```
methods(class = class(dat)[3])
```

```
[[<-
                                                                   $
##
    [1] [
                       [<-
    [6] $<-
                       aggregate
                                      anyDuplicated as.data.frame as.list
## [11] as.matrix
                                      cbind
                                                     coerce
                                                                   dim
  [16] dimnames
                       dimnames<-
                                      droplevels
                                                     duplicated
                                                                   edit
## [21] format
                       formula
                                      head
                                                     initialize
                                                                   is.na
```

```
## [26] Math
                                        na.exclude
                                                       na.omit
                                                                       Ops
                        merge
## [31] plot
                                                       rbind
                                                                       row.names
                        print
                                        prompt
## [36] row.names<-
                        rowsum
                                        show
                                                       slotsFromS3
                                                                       split
## [41] split<-
                        stack
                                        str
                                                       subset
                                                                       summary
## [46] Summary
                                        tail
                                                       transform
                                                                       unique
## [51] unstack
                        within
## see '?methods' for accessing help and source code
  3. If you call print(dat), what print method is being dispatched? (1 point)
# print(dat)
# print.default(dat)
The default print method is dispatched.
  4. Set the class of dat to be a data frame. (1 point)
class(dat) <- "data.frame"</pre>
  5. If you call print(dat) again, what print method is being dispatched? (1 point)
# print(dat)
The data.frame print method is dispatched.
Define a new generic function nUnique with the code below.
nUnique <- function(x) {
    UseMethod('nUnique')
}
  6. Write a default method for nUnique to count the number of unique values in an element. (2 points)
nUnique.default <- function(x){
  return(length(unique(x)))
  7. Check your function (2 points)
nUnique(letters) # should return 26
## [1] 26
nUnique(sample(10, 100, replace = TRUE)) # should return 10 (probably)
## [1] 10
  8. Write a data frame method for nUnique to operate on data frame objects. This version should return
     counts for each column in a data.frame. (2 points)
nUnique.data.frame <- function(x){</pre>
  z <- sapply(1:ncol(x), function(i) length(unique(x[,i])))</pre>
  return(z)
}
  9. Check your function (2 points)
nUnique(dat)
```

[1] 39 8321 8320 15

Question 2

15 points

##

Programming with classes. The following function will generate random patient information.

```
makePatient <- function() {
  vowel <- grep("[aeiou]", letters)
  cons <- grep("[^aeiou]", letters)
  name <- paste(sample(LETTERS[cons], 1), sample(letters[vowel], 1), sample(letters[cons], 1), sep='')
  gender <- factor(sample(0:1, 1), levels=0:1, labels=c('female', 'male'))
  dob <- as.Date(sample(7500, 1), origin="1970-01-01")
  n <- sample(6, 1)
  doa <- as.Date(sample(1500, n), origin="2010-01-01")
  pulse <- round(rnorm(n, 80, 10))
  temp <- round(rnorm(n, 98.4, 0.3), 2)
  fluid <- round(runif(n), 2)
  list(name, gender, dob, doa, pulse, temp, fluid)
}</pre>
```

1. Create an S3 class medicalRecord for objects that are a list with the named elements name, gender, date_of_birth, date_of_admission, pulse, temperature, fluid_intake. Note that an individual patient may have multiple measurements for some measurements. Set the RNG seed to 8 and create a medical record by taking the output of makePatient. Print the medical record, and print the class of the medical record. (5 points)

```
mrf <- function(){</pre>
  set.seed(8)
  mr <- makePatient()</pre>
  names(mr)<-c("name", "gender", "date_of_birth", "date_of_admission", "pulse", "temperature", "fluid_int</pre>
  class(mr) <- "medicalRecord"</pre>
  return(mr)
}
mr <- mrf()
mr
## $name
## [1] "Mev"
##
## $gender
## [1] male
## Levels: female male
##
## $date_of_birth
## [1] "1976-08-09"
## $date_of_admission
## [1] "2011-03-14" "2013-10-30" "2013-02-27" "2012-08-23" "2011-11-16"
##
## $pulse
## [1] 67 81 95 74 81
## $temperature
## [1] 98.33 98.16 99.00 98.49 98.67
```

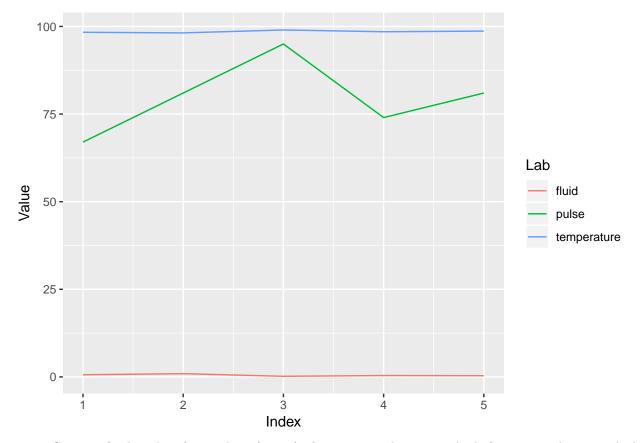
```
## $fluid intake
## [1] 0.62 0.93 0.18 0.39 0.34
## attr(,"class")
## [1] "medicalRecord"
  2. Write a medicalRecord method for the generic function mean, which returns averages for pulse,
     temperature and fluids. Also write a medical Record method for print, which employs some nice
     formatting, perhaps arranging measurements by date, and plot, that generates a composite plot of
     measurements over time. Call each function for the medical record created in part 1. (5 points)
# mean function
mean.medicalRecord <- function(x){</pre>
  cat(sprintf("Average for pulse: %f\nAverage for temperature: %f\nAverage for fluids: %f", mean(x$puls
mean(mr)
## Average for pulse: 79.600000
## Average for temperature: 98.530000
## Average for fluids: 0.492000
# print function
print.medicalRecord <- function(x) {</pre>
  date_admission <- x$date_of_admission[order(x$date_of_admission)]</pre>
  pulse <- x$pulse[order(x$date_of_admission)]</pre>
 temperature <- x$temperature[order(x$date_of_admission)]</pre>
 fluid <- x$fluid_intake[order(x$date_of_admission)]</pre>
  cat(sprintf("name: %s\ngender: %d\ndate of birth: %s\ndate of admission: %s\npulse: %d\ntemperature:
}
mr
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2011-03-14
## pulse: 67
## temperature: 98.330000
## fluid intake: 0.620000
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2011-11-16
## pulse: 81
## temperature: 98.670000
## fluid intake: 0.340000
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2012-08-23
## pulse: 74
## temperature: 98.490000
```

fluid intake: 0.390000

name: Mev

```
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2013-02-27
## pulse: 95
## temperature: 99.000000
## fluid intake: 0.180000
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2013-10-30
## pulse: 81
## temperature: 98.160000
## fluid intake: 0.930000
##
# plot function
plot.medicalRecord <- function(x){</pre>
  pulse <- x$pulse
 temperature <- x$temperature
  fluid <- x$fluid_intake</pre>
  Value <- c(pulse, temperature, fluid)</pre>
  Index <- 1:length(pulse)</pre>
  Lab <- c(rep("pulse",length(pulse)),rep("temperature",length(temperature)),rep("fluid",length(fluid))
  df <- data.frame(Index, Value, Lab)</pre>
  library(ggplot2)
  ggplot(data=df,aes(x=Index,y=Value,group=Lab)) + geom_line(aes(color=Lab))
}
plot(mr)
```

Warning: package 'ggplot2' was built under R version 3.4.4



3. Create a further class for a cohort (group) of patients, and write methods for mean and print which, when applied to a cohort, apply mean or print to each patient contained in the cohort. Hint: think of this as a "container" for patients. Reset the RNG seed to 8 and create a cohort of ten patients, then show the output for mean and print. (5 points)

```
set.seed(8)
patients <- replicate(10,list(makePatient()))</pre>
class(patients) <- "Patients"</pre>
# mean
mean.Patients <- function(x){</pre>
  pulse <- sapply(1:10, function(i) mean(x[[i]][[5]]))</pre>
  temperature <- sapply(1:10, function(i) mean(x[[i]][[6]]))</pre>
  fluid <- sapply(1:10, function(i) mean(x[[i]][[7]]))</pre>
  message_pulse <- sprintf("Mean pulse for the subject %d: %f",1:10,pulse)
  message_temp <- sprintf("Mean temperature for the subject %d: %f",1:10,temperature)
  message_fluid <- sprintf("Mean fluid for the subject %d: %f",1:10,fluid)
  return(c(message_pulse,message_temp, message_fluid))
}
mean(patients)
    [1] "Mean pulse for the subject 1: 79.600000"
##
```

[2] "Mean pulse for the subject 2: 78.000000"

[3] "Mean pulse for the subject 3: 81.500000" [4] "Mean pulse for the subject 4: 78.000000"

##

##

```
## [5] "Mean pulse for the subject 5: 88.333333"
## [6] "Mean pulse for the subject 6: 83.500000"
## [7] "Mean pulse for the subject 7: 83.000000"
## [8] "Mean pulse for the subject 8: 77.500000"
## [9] "Mean pulse for the subject 9: 77.000000"
## [10] "Mean pulse for the subject 10: 79.333333"
## [11] "Mean temperature for the subject 1: 98.530000"
## [12] "Mean temperature for the subject 2: 98.495000"
## [13] "Mean temperature for the subject 3: 98.440000"
## [14] "Mean temperature for the subject 4: 98.600000"
## [15] "Mean temperature for the subject 5: 98.050000"
## [16] "Mean temperature for the subject 6: 98.450000"
## [17] "Mean temperature for the subject 7: 98.010000"
## [18] "Mean temperature for the subject 8: 98.148333"
## [19] "Mean temperature for the subject 9: 98.830000"
## [20] "Mean temperature for the subject 10: 98.300000"
## [21] "Mean fluid for the subject 1: 0.492000"
## [22] "Mean fluid for the subject 2: 0.245000"
## [23] "Mean fluid for the subject 3: 0.403333"
## [24] "Mean fluid for the subject 4: 0.650000"
## [25] "Mean fluid for the subject 5: 0.586667"
## [26] "Mean fluid for the subject 6: 0.452500"
## [27] "Mean fluid for the subject 7: 0.970000"
## [28] "Mean fluid for the subject 8: 0.336667"
## [29] "Mean fluid for the subject 9: 0.445000"
## [30] "Mean fluid for the subject 10: 0.658333"
# print
print.Patients <- function(x){</pre>
  # get the number of visits from each person
  n_visit <- sapply(1:10, function(i) length(x[[i]][[4]]))</pre>
  # store the admission date info
  date_admission <- sapply(1:10, function(i) x[[i]][[4]][order(x[[i]][[4]])])
  # pulse
  pulse <- sapply(1:10, function(i) x[[i]][[5]][order(x[[i]][[4]])])</pre>
  temperature \leftarrow sapply(1:10, function(i) x[[i]][[6]][order(x[[i]][[4]])])
  fluid <- sapply(1:10, function(i) x[[i]][[7]][order(x[[i]][[4]])])
  # print
  sapply(1:10, function(i) cat(sprintf("name: %s\ngender: %d\ndate of birth: %s\ndate of admission: %s\
}
patients
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2011-03-14
## pulse: 67
## temperature: 98.330000
```

```
## fluid intake: 0.620000
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2011-11-16
## pulse: 81
## temperature: 98.670000
## fluid intake: 0.340000
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2012-08-23
## pulse: 74
## temperature: 98.490000
## fluid intake: 0.390000
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2013-02-27
## pulse: 95
## temperature: 99.000000
## fluid intake: 0.180000
## name: Mev
## gender: 2
## date of birth: 1976-08-09
## date of admission: 2013-10-30
## pulse: 81
## temperature: 98.160000
## fluid intake: 0.930000
##
## name: Yul
## gender: 2
## date of birth: 1988-06-28
## date of admission: 2012-01-16
## pulse: 76
## temperature: 98.920000
## fluid intake: 0.140000
## name: Yul
## gender: 2
## date of birth: 1988-06-28
## date of admission: 2013-08-07
## pulse: 80
## temperature: 98.070000
## fluid intake: 0.350000
##
## name: Zet
## gender: 1
## date of birth: 1970-06-13
## date of admission: 2010-03-21
## pulse: 79
## temperature: 98.580000
## fluid intake: 0.220000
```

name: Zet
gender: 1

```
## date of birth: 1970-06-13
## date of admission: 2010-04-01
```

pulse: 73

temperature: 98.320000 ## fluid intake: 0.610000

name: Zet
gender: 1

date of birth: 1970-06-13 ## date of admission: 2012-08-29

pulse: 88

temperature: 98.470000 ## fluid intake: 0.590000

name: Zet
gender: 1

date of birth: 1970-06-13 ## date of admission: 2013-06-01

pulse: 84

temperature: 98.220000 ## fluid intake: 0.250000

name: Zet
gender: 1

date of birth: 1970-06-13 ## date of admission: 2013-11-03

pulse: 72

temperature: 98.540000 ## fluid intake: 0.030000

name: Zet
gender: 1

date of birth: 1970-06-13 ## date of admission: 2014-02-05

pulse: 93

temperature: 98.510000
fluid intake: 0.720000

##

name: Qih
gender: 1

date of birth: 1987-08-30 ## date of admission: 2011-06-22

pulse: 78

temperature: 98.600000 ## fluid intake: 0.650000

##

name: Wut
gender: 2

date of birth: 1974-06-28 ## date of admission: 2010-04-12

pulse: 76

temperature: 98.050000 ## fluid intake: 0.650000

name: Wut
gender: 2

date of birth: 1974-06-28 ## date of admission: 2011-02-16

pulse: 93

```
## temperature: 98.260000
## fluid intake: 0.970000
## name: Wut
## gender: 2
## date of birth: 1974-06-28
## date of admission: 2012-04-12
## pulse: 96
## temperature: 97.840000
## fluid intake: 0.140000
##
## name: Juy
## gender: 2
## date of birth: 1983-06-09
## date of admission: 2010-03-10
## pulse: 81
## temperature: 99.110000
## fluid intake: 0.660000
## name: Juy
## gender: 2
## date of birth: 1983-06-09
## date of admission: 2010-03-25
## pulse: 90
## temperature: 98.580000
## fluid intake: 0.260000
## name: Juy
## gender: 2
## date of birth: 1983-06-09
## date of admission: 2010-04-18
## pulse: 75
## temperature: 98.580000
## fluid intake: 0.600000
## name: Juy
## gender: 2
## date of birth: 1983-06-09
## date of admission: 2010-06-10
## pulse: 88
## temperature: 97.530000
## fluid intake: 0.290000
##
## name: God
## gender: 1
## date of birth: 1990-02-12
## date of admission: 2010-03-12
## pulse: 83
## temperature: 98.010000
## fluid intake: 0.970000
## name: Fut
## gender: 2
## date of birth: 1970-01-11
## date of admission: 2011-04-07
## pulse: 80
```

temperature: 97.870000 ## fluid intake: 0.360000

```
## name: Fut
## gender: 2
## date of birth: 1970-01-11
## date of admission: 2011-04-14
## pulse: 83
## temperature: 97.910000
## fluid intake: 0.000000
## name: Fut
## gender: 2
## date of birth: 1970-01-11
## date of admission: 2011-08-16
## pulse: 66
## temperature: 98.490000
## fluid intake: 0.130000
## name: Fut
## gender: 2
## date of birth: 1970-01-11
## date of admission: 2013-03-15
## pulse: 74
## temperature: 98.380000
## fluid intake: 0.310000
## name: Fut
## gender: 2
## date of birth: 1970-01-11
## date of admission: 2013-06-20
## pulse: 74
## temperature: 98.410000
## fluid intake: 0.490000
## name: Fut
## gender: 2
## date of birth: 1970-01-11
## date of admission: 2013-11-12
## pulse: 88
## temperature: 97.830000
## fluid intake: 0.730000
## name: Pet
## gender: 2
## date of birth: 1979-01-01
## date of admission: 2010-10-30
## pulse: 85
## temperature: 98.840000
## fluid intake: 0.600000
## name: Pet
## gender: 2
## date of birth: 1979-01-01
## date of admission: 2012-05-10
## pulse: 69
## temperature: 98.820000
## fluid intake: 0.290000
##
## name: Yed
## gender: 2
```

date of birth: 1977-11-11

```
## date of admission: 2010-01-28
```

pulse: 63

temperature: 97.950000
fluid intake: 0.940000

name: Yed
gender: 2

date of birth: 1977-11-11 ## date of admission: 2010-03-06

pulse: 81

temperature: 98.450000 ## fluid intake: 0.670000

name: Yed
gender: 2

date of birth: 1977-11-11 ## date of admission: 2010-07-10

pulse: 98

temperature: 98.650000 ## fluid intake: 0.790000

name: Yed
gender: 2

date of birth: 1977-11-11 ## date of admission: 2010-08-27

pulse: 66

temperature: 97.680000 ## fluid intake: 0.360000

name: Yed
gender: 2

date of birth: 1977-11-11 ## date of admission: 2011-06-18

pulse: 83

temperature: 98.000000 ## fluid intake: 0.690000

name: Yed
gender: 2

date of birth: 1977-11-11 ## date of admission: 2013-01-06

pulse: 85

temperature: 99.070000 ## fluid intake: 0.500000

##