

BIOS 6301: Assignment 8

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Due Tuesday, 14 November, 1:00 PM

$5^{n=\text{day}}$ 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

```
library(readxl)
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, mode = 'wb')
}
dat <- readxl::read_excel(fn, sheet = 2)
```

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

Object 'dat' has 3 classes: 'tbl_df', 'tbl', and 'data.frame'.

```
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="tbl_df")
```

```
## [1] [          [[          [[<-          [<-          $
## [6] $<-        as.data.frame coerce          initialize names<-
## [11] Ops        row.names<- show          slotsFromS3 str
## see '?methods' for accessing help and source code
```

```
methods(class="tbl")
```

```
## [1] [[<-      [<-      $<-      coerce      format      initialize
## [7] Ops        print      show      slotsFromS3
## see '?methods' for accessing help and source code
```

```
methods(class="data.frame")
```

```
## [1] [          [[          [[<-          [<-          $<-
## [6] aggregate anyDuplicated anyNA      as.data.frame as.list
## [11] as.matrix  as.vector  by      cbind        coerce
## [16] dim        dimnames   dimnames<- droplevels   duplicated
## [21] edit       format     formula   head         initialize
## [26] is.na      Math       merge     na.exclude   na.omit
## [31] Ops        plot       print     prompt       rbind
## [36] row.names  row.names<- rowsum    show         slotsFromS3
## [41] split      split<-    stack     str          subset
## [46] summary    Summary    t         tail         transform
## [51] type.convert unique      unstack   within       xtfm
## see '?methods' for accessing help and source code
```

3. If you call `print(dat)`, what print method is being dispatched? (1 point)

There is a 'print.tbl_df' method within print. Since we know 'tbl_df' is the first class type for 'dat', then 'print.tbl_df' is the method that is being dispatched when we call `print(dat)`.

```
#methods(print)
print(head(dat))
```

```
## # A tibble: 6 x 4
##   Procedure Code Categ~1 'ICD-10-PCS Codes' Procedure Code Descr~2 'Code Status'
##   <chr>                  <chr>                  <chr>                  <chr>
## 1 AAA                  04B00ZZ                Excision of Abdominal~ No change
## 2 AAA                  04B04ZZ                Excision of Abdominal~ No change
## 3 AAA                  04R007Z                Replacement of Abdomi~ No change
## 4 AAA                  04R00JZ                Replacement of Abdomi~ No change
## 5 AAA                  04R00KZ                Replacement of Abdomi~ No change
## 6 AAA                  04R047Z                Replacement of Abdomi~ No change
## # i abbreviated names: 1: 'Procedure Code Category',
## #   2: 'Procedure Code Descriptions'
```

4. Set the class of `dat` to be a `data.frame`. (1 point)

```
dat <- as.data.frame(dat)
class(dat)
```

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

5. If you call `print(dat)` again, what print method is being dispatched? (1 point)

Since 'dat' is now a `data.frame`, when we call the print command, the `print.data.frame` method is being dispatched

```
print(head(dat))
```

```
## Procedure Code Category ICD-10-PCS Codes
## 1 AAA 04B00ZZ
## 2 AAA 04B04ZZ
## 3 AAA 04R007Z
## 4 AAA 04R00JZ
## 5 AAA 04R00KZ
## 6 AAA 04R047Z
##
## Procedure Code Descriptions
## 1 Excision of Abdominal Aorta, Open Approach
## 2 Excision of Abdominal Aorta, Percutaneous Endoscopic Approach
## 3 Replacement of Abdominal Aorta with Autologous Tissue Substitute, Open Approach
## 4 Replacement of Abdominal Aorta with Synthetic Substitute, Open Approach
## 5 Replacement of Abdominal Aorta with Nonautologous Tissue Substitute, Open Approach
## 6 Replacement of Abdominal Aorta with Autologous Tissue Substitute, Percutaneous Endoscopic Approach
## Code Status
## 1 No change
## 2 No change
## 3 No change
## 4 No change
## 5 No change
## 6 No change
```

Define a new generic function `nUnique` with the code below.

```
nUnique <- function(x) {
  UseMethod('nUnique')
}
```

- Write a default method for `nUnique` to count the number of unique values in an element. (2 points)

Using the `nUnique` default method, I was able to determine the number of unique values for two variables within the 'dat' dataset. For 'Procedure Code Category' I found 39 unique values, and for 'ICD-10-PCS Codes' I found 8993 unique values.

```
nUnique.default <- function(x) {
  length(unique(x))
}

v1 <- dat$`Procedure Code Category`
nUnique(v1)
```

```
## [1] 39
```

```
v2 <- dat$`ICD-10-PCS Codes`
nUnique(v2)
```

```
## [1] 8993
```

- Check your function (2 points)

```
nUnique(letters) # should return 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) {  
  for(i in colnames(x)){  
    cat("Unique values in", i, ":", length(unique(x[,i])), "\n")  
  }  
}
```

9. Check your function (2 points)

```
nUnique(dat)
```

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)  
}
```

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)

```
set.seed(8)  
p <- makePatient()  
names(p) <- c("name", "gender", "date_of_birth", "date_of_admission", "pulse", "temperature", "fluid_intake")  
class(p) <- 'medicalRecord'  
print(p)
```

```
## $name
## [1] "Yes"
##
## $gender
## [1] male
## Levels: female male
##
## $date_of_birth
## [1] "1977-05-03"
##
## $date_of_admission
## [1] "2013-06-09" "2013-07-02"
##
## $pulse
## [1] 79 78
##
## $temperature
## [1] 98.07 97.50
##
## $fluid_intake
## [1] 0.28 0.52
##
## 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 `medicalRecord` 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.medicalRecord <- function(x) {
  list(mean(p[[5]]), mean(p[[6]]), mean(p[[7]]))
}
```

```
p2 <- mean(p)
names(p2) <- c("Avg. Pulse", "Avg. Temperature", " Avg. Fluid Intake")
print(p2)
```

```
## $'Avg. Pulse'
## [1] 78.5
##
## $'Avg. Temperature'
## [1] 97.785
##
## $' Avg. Fluid Intake'
## [1] 0.4
```

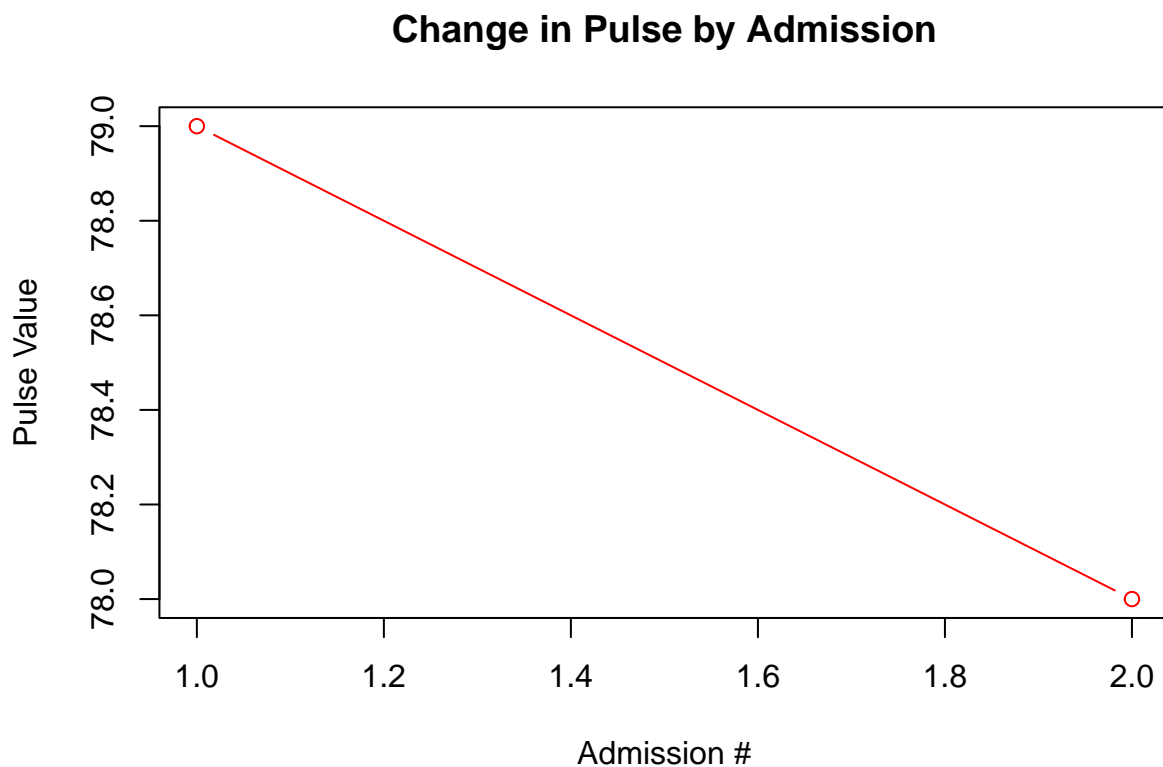
```
print.medicalRecord<- function(med) {
  cat(sprintf("Date of Admission: %s\nName: %s\nGender: %s\nDate of Birth: %s\nPulse: %s\nTemperature: %s\nFluid Intake: %s",
    med$date_of_admission, med$name, med$gender, med$date_of_birth, med$pulse, med$temperature, med$fluid_intake))
}

print(p)
```

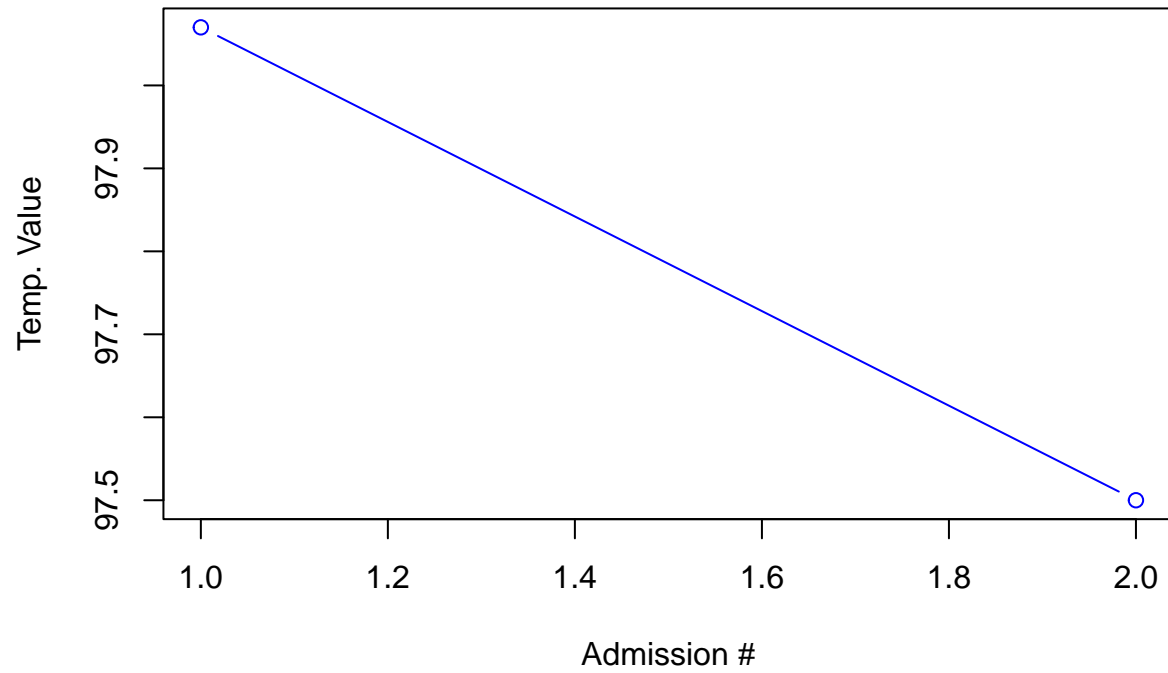
```
## Date of Admission: 2013-06-09
## Name: Yes
## Gender: male
## Date of Birth: 1977-05-03
## Pulse: 79
## Temperature: 98.07
## Fluid Intake 0.28
## Date of Admission: 2013-07-02
## Name: Yes
## Gender: male
## Date of Birth: 1977-05-03
## Pulse: 78
## Temperature: 97.5
## Fluid Intake 0.52
##
```

```
plot.medicalRecord <- function(x){
  plot(x[[5]],type='b',col='red',main="Change in Pulse by Admission",xlab='Admission #', ylab='Pulse Value')
  plot(x[[6]],type='b',col='blue',main="Change in Temperature by Admission",xlab='Admission #', ylab='Temperature')
  plot(x[[7]],type='b',col='green',main="Change in Fluid Intake by Admission",xlab='Admission #', ylab='Fluid Intake')
}

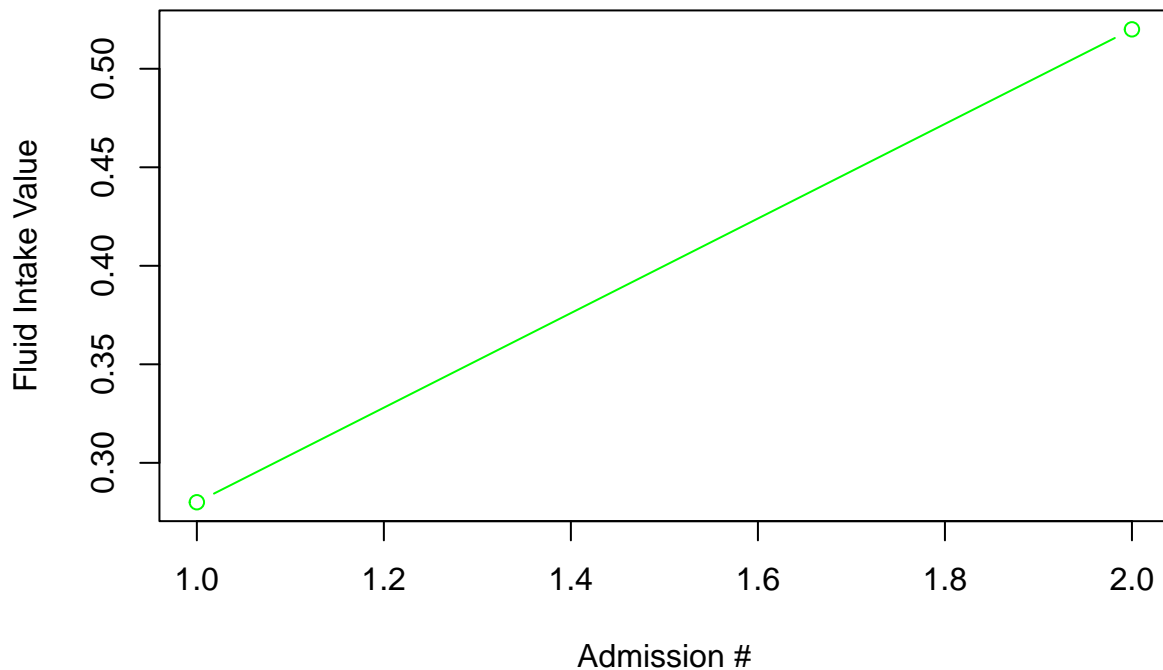
plot(p)
```



Change in Temperature by Admission



Change in Fluid Intake by Admission



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)

```
rm(.Random.seed, envir = globalenv())
set.seed(8)
cohort = list()
for (i in 1:10){
  x = makePatient()
  names(x) <- c("name", "gender", "date_of_birth", "date_of_admission", "pulse", "temperature", "fluid_intake")
  class(x) <- 'medicalRecord'
  cohort[[i]] = x
}
class(cohort) <- 'cohort'
class(cohort)
```

```
## [1] "cohort"
```

```
#Mean Method for Cohort
mean.cohort <- function(x){
  lapply(x, mean)
}

mean(cohort)
```



```

## [[1]]
## [[1]][[1]]
## [1] 78.5
##
## [[1]][[2]]
## [1] 97.785
##
## [[1]][[3]]
## [1] 0.4
##
##
## [[2]]
## [[2]][[1]]
## [1] 78.5
##
## [[2]][[2]]
## [1] 97.785
##
## [[2]][[3]]
## [1] 0.4
##
##
## [[3]]
## [[3]][[1]]
## [1] 78.5
##
## [[3]][[2]]
## [1] 97.785
##
## [[3]][[3]]
## [1] 0.4
##
##
## [[4]]
## [[4]][[1]]
## [1] 78.5
##
## [[4]][[2]]
## [1] 97.785
##
## [[4]][[3]]
## [1] 0.4
##
##
## [[5]]
## [[5]][[1]]
## [1] 78.5
##
## [[5]][[2]]
## [1] 97.785
##
## [[5]][[3]]
## [1] 0.4
##

```

```

##
## [[6]]
## [[6]][[1]]
## [1] 78.5
##
## [[6]][[2]]
## [1] 97.785
##
## [[6]][[3]]
## [1] 0.4
##
##
## [[7]]
## [[7]][[1]]
## [1] 78.5
##
## [[7]][[2]]
## [1] 97.785
##
## [[7]][[3]]
## [1] 0.4
##
##
## [[8]]
## [[8]][[1]]
## [1] 78.5
##
## [[8]][[2]]
## [1] 97.785
##
## [[8]][[3]]
## [1] 0.4
##
##
## [[9]]
## [[9]][[1]]
## [1] 78.5
##
## [[9]][[2]]
## [1] 97.785
##
## [[9]][[3]]
## [1] 0.4
##
##
## [[10]]
## [[10]][[1]]
## [1] 78.5
##
## [[10]][[2]]
## [1] 97.785
##
## [[10]][[3]]
## [1] 0.4

```

```

#Print Method for Cohort
print.cohort <- function(x){
  for (i in 1:10){
    print(cohort[[i]])
  }
}

print(cohort)

```

```

## Date of Admission: 2013-06-09
## Name: Yes
## Gender: male
## Date of Birth: 1977-05-03
## Pulse: 79
## Temperature: 98.07
## Fluid Intake 0.28
## Date of Admission: 2013-07-02
## Name: Yes
## Gender: male
## Date of Birth: 1977-05-03
## Pulse: 78
## Temperature: 97.5
## Fluid Intake 0.52
##
## Date of Admission: 2010-11-16
## Name: Fal
## Gender: male
## Date of Birth: 1988-05-24
## Pulse: 76
## Temperature: 98.23
## Fluid Intake 0.18
## Date of Admission: 2013-09-12
## Name: Fal
## Gender: male
## Date of Birth: 1988-05-24
## Pulse: 96
## Temperature: 98.75
## Fluid Intake 0.96
## Date of Admission: 2013-03-24
## Name: Fal
## Gender: male
## Date of Birth: 1988-05-24
## Pulse: 87
## Temperature: 98.21
## Fluid Intake 0.1
##
## Date of Admission: 2013-03-25
## Name: Zog
## Gender: male
## Date of Birth: 1988-12-14
## Pulse: 69
## Temperature: 98.49
## Fluid Intake 0.81

```

Date of Admission: 2013-07-29
Name: Zog
Gender: male
Date of Birth: 1988-12-14
Pulse: 75
Temperature: 98.82
Fluid Intake 0.59
Date of Admission: 2013-10-27
Name: Zog
Gender: male
Date of Birth: 1988-12-14
Pulse: 80
Temperature: 98.74
Fluid Intake 0.28
Date of Admission: 2010-02-24
Name: Zog
Gender: male
Date of Birth: 1988-12-14
Pulse: 84
Temperature: 98.54
Fluid Intake 0.4

Date of Admission: 2014-01-28
Name: Yol
Gender: male
Date of Birth: 1986-03-11
Pulse: 69
Temperature: 98.29
Fluid Intake 0.03
Date of Admission: 2013-03-24
Name: Yol
Gender: male
Date of Birth: 1986-03-11
Pulse: 78
Temperature: 98.44
Fluid Intake 0.13
Date of Admission: 2012-03-10
Name: Yol
Gender: male
Date of Birth: 1986-03-11
Pulse: 87
Temperature: 98.78
Fluid Intake 0.12
Date of Admission: 2010-02-22
Name: Yol
Gender: male
Date of Birth: 1986-03-11
Pulse: 84
Temperature: 98.87
Fluid Intake 0.39
Date of Admission: 2011-12-27
Name: Yol
Gender: male
Date of Birth: 1986-03-11

```

## Pulse: 89
## Temperature: 98.27
## Fluid Intake 0.97
## Date of Admission: 2012-11-26
## Name: Yol
## Gender: male
## Date of Birth: 1986-03-11
## Pulse: 92
## Temperature: 98.26
## Fluid Intake 0.14
##
## Date of Admission: 2012-08-30
## Name: Yak
## Gender: female
## Date of Birth: 1983-09-15
## Pulse: 90
## Temperature: 98.58
## Fluid Intake 0.26
## Date of Admission: 2012-04-07
## Name: Yak
## Gender: female
## Date of Birth: 1983-09-15
## Pulse: 88
## Temperature: 97.53
## Fluid Intake 0.29
## Date of Admission: 2011-07-19
## Name: Yak
## Gender: female
## Date of Birth: 1983-09-15
## Pulse: 75
## Temperature: 98.58
## Fluid Intake 0.6
## Date of Admission: 2012-07-11
## Name: Yak
## Gender: female
## Date of Birth: 1983-09-15
## Pulse: 81
## Temperature: 99.11
## Fluid Intake 0.66
##
## Date of Admission: 2012-04-24
## Name: Gaf
## Gender: female
## Date of Birth: 1978-04-27
## Pulse: 89
## Temperature: 98.32
## Fluid Intake 0.42
## Date of Admission: 2010-07-19
## Name: Gaf
## Gender: female
## Date of Birth: 1978-04-27
## Pulse: 91
## Temperature: 98.01
## Fluid Intake 0.47

```

Date of Admission: 2012-08-06
Name: Gaf
Gender: female
Date of Birth: 1978-04-27
Pulse: 77
Temperature: 98.96
Fluid Intake 0.74
Date of Admission: 2013-08-21
Name: Gaf
Gender: female
Date of Birth: 1978-04-27
Pulse: 75
Temperature: 98.52
Fluid Intake 0.62
Date of Admission: 2011-05-03
Name: Gaf
Gender: female
Date of Birth: 1978-04-27
Pulse: 90
Temperature: 98.61
Fluid Intake 0.36

Date of Admission: 2011-09-16
Name: Kuw
Gender: female
Date of Birth: 1980-11-07
Pulse: 72
Temperature: 98.21
Fluid Intake 0.29
Date of Admission: 2010-10-29
Name: Kuw
Gender: female
Date of Birth: 1980-11-07
Pulse: 81
Temperature: 98.17
Fluid Intake 0.93
Date of Admission: 2012-07-10
Name: Kuw
Gender: female
Date of Birth: 1980-11-07
Pulse: 71
Temperature: 98.65
Fluid Intake 0.25
Date of Admission: 2010-10-03
Name: Kuw
Gender: female
Date of Birth: 1980-11-07
Pulse: 82
Temperature: 98.49
Fluid Intake 0.12

Date of Admission: 2012-03-02
Name: Mav
Gender: female

Date of Birth: 1989-07-16
Pulse: 63
Temperature: 99.07
Fluid Intake 0.01
Date of Admission: 2010-06-11
Name: Mav
Gender: female
Date of Birth: 1989-07-16
Pulse: 83
Temperature: 98.45
Fluid Intake 0.79
Date of Admission: 2010-02-08
Name: Mav
Gender: female
Date of Birth: 1989-07-16
Pulse: 66
Temperature: 97.95
Fluid Intake 0.79
Date of Admission: 2010-04-19
Name: Mav
Gender: female
Date of Birth: 1989-07-16
Pulse: 88
Temperature: 98
Fluid Intake 0.5

Date of Admission: 2012-06-24
Name: Fel
Gender: male
Date of Birth: 1985-08-16
Pulse: 65
Temperature: 98.21
Fluid Intake 0.06
Date of Admission: 2010-09-26
Name: Fel
Gender: male
Date of Birth: 1985-08-16
Pulse: 81
Temperature: 98.51
Fluid Intake 0.24

Date of Admission: 2010-03-14
Name: Say
Gender: female
Date of Birth: 1974-09-22
Pulse: 77
Temperature: 98.54
Fluid Intake 0.15
##