

Bios 6301: Assignment 5

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Due Tuesday, 15 November, 1:00 PM

$5^{n=\text{day}}$ points taken off for each day late.

50 points total.

Submit a single knitr file (named `homework5.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 `homework5.rmd` or include author name may result in 5 points taken off.

Question 1

24 points

Import the HAART dataset (`haart.csv`) from the GitHub repository into R, and perform the following manipulations: (4 points each)

```
haart <- read.csv("https://raw.githubusercontent.com/fonnesbeck/Bios6301/master/datasets/haart.csv",
  stringsAsFactors = FALSE)
```

```
##### 1. Convert date columns into a usable (for analysis) format. Use the
##### `table` command to display the counts of the year from `init.date`.
```

```
library(lubridate)
```

```
## Warning: package 'lubridate' was built under R version 3.2.5
```

```
##
```

```
## Attaching package: 'lubridate'
```

```
## The following object is masked from 'package:base':
```

```
##
```

```
##      date
```

```
haart[, "init.date"] <- mdy(haart[, "init.date"])
haart[, "last.visit"] <- mdy(haart[, "last.visit"])
haart[, "date.death"] <- mdy(haart[, "date.death"])
```

```
years <- substr(haart[, "init.date"], 1, 4)
table(years)
```

```
## years
```

```
## 1998 2000 2001 2002 2003 2004 2005 2006 2007
```

```
##    1    5   17   60  270  292  207  104   44
```

```
##### 2. Create an indicator variable (one which takes the values 0 or 1 only)
##### to represent death within 1 year of the initial visit. How many
##### observations died in year 1?
```

```
haart[, "death.1yr"] <- 0
ind <- which(haart[, "death"] == 1)
time <- as.numeric(haart[ind, "date.death"] - haart[ind, "init.date"])
temp <- data.frame(ind, time)
haart[temp[, "ind"], "death.1yr"] <- as.numeric(temp[, "time"] <= 365)

length(which(haart[, "death.1yr"] == 1))
```

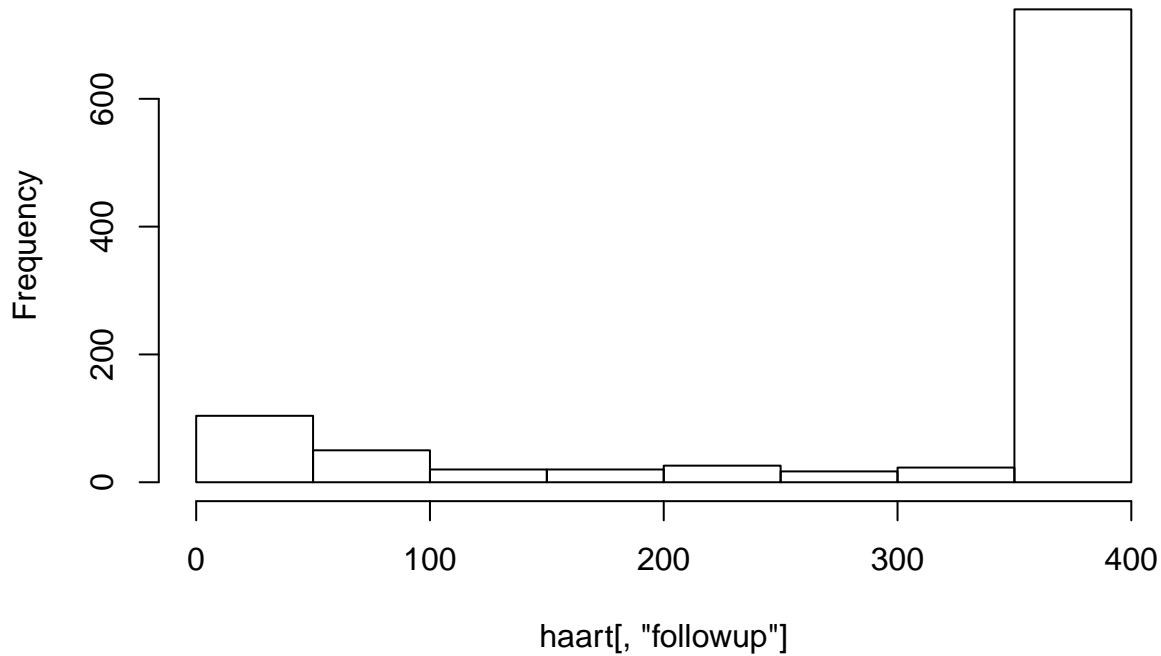
```
## [1] 92
```

```
##### 3. Use the `init.date`, `last.visit` and `death.date` columns to calculate
##### a followup time (in days), which is the difference between the first and
##### either the last visit or a death event (whichever comes first). If these
##### times are longer than 1 year, censor them (this means if the value is
##### above 365, set followup to 365). Print the quantile for this new
##### variable.
```

```
haart[ind, "followup"] <- temp[, "time"]
ind.2 <- which(haart[, "death"] == 0)
time.2 <- as.numeric(haart[ind.2, "last.visit"] - haart[ind.2, "init.date"])
temp.2 <- data.frame(ind.2, time.2)
haart[ind.2, "followup"] <- temp.2[, "time.2"]
# hist(haart[, 'followup']) quantile(haart[, 'followup'])

for (i in 1:nrow(haart)) {
  if (haart[i, "followup"] >= 365) {
    haart[i, "followup"] = 365
  }
}
hist(haart[, "followup"])
```

Histogram of haart[, "followup"]



```
quantile(haart[, "followup"])
```

```
##      0%    25%    50%    75%   100%
##      0.0  329.5  365.0  365.0  365.0
```

```
##### 4. Create another indicator variable representing loss to followup; this
##### means the observation is not known to be dead but does not have any
##### followup visits after the first year. How many records are
##### lost-to-followup?
```

```
for (i in 1:nrow(haart)) {
  if (haart[i, "followup"] < 365 & haart[i, "death"] == 0) {
    haart[i, "lossfu"] = 1
  } else {
    haart[i, "lossfu"] = 0
  }
}
length(which(haart[, "lossfu"] == 1))
```

```
## [1] 173
```

```
##### 5. Recall our work in class, which separated the `init.reg` field into a
##### set of indicator variables, one for each unique drug. Create these fields
##### and append them to the database as new columns. Which drug regimen are
##### found over 100 times?
```

```
all.reg <- unique(unlist(strsplit(haart[, "init.reg"], ",")))
all.reg
```

```
## [1] "3TC" "AZT" "EFV" "NVP" "D4T" "ABC" "DDI" "IDV" "LPV" "RTV" "SQV"
## [12] "FTC" "TDF" "DDC" "NFV" "T20" "ATV" "FPV"
```

```
haart[, all.reg] <- 0

for (i in 1:nrow(haart)) {
  reg <- unlist(strsplit(haart[i, "init.reg"], ","))
  haart[i, reg] <- 1
}

regimen <- haart[, "init.reg"]
z <- as.data.frame(table(regimen))
z[which(z[, "Freq"] > 100), ]
```

```
##           regimen Freq
## 10 3TC,AZT,EFV  421
## 17 3TC,AZT,NVP  284
```

6. The dataset `haart2.csv` contains a few additional observations for the same study. Import these and append them to your master dataset (if you were smart about how you coded the previous steps, cleaning the additional observations should be easy!). Show the first five records and the last five records of the complete (and clean) data set.

```
haart2 <- read.csv("https://raw.githubusercontent.com/fonnesbeck/Bios6301/master/datasets/haart2.csv",
  stringsAsFactors = FALSE)

haart2[, "init.date"] <- mdy(haart2[, "init.date"])
haart2[, "last.visit"] <- mdy(haart2[, "last.visit"])
haart2[, "date.death"] <- mdy(haart2[, "date.death"])
```

```
## Warning: All formats failed to parse. No formats found.
```

```
haart2[, "death.1yr"] <- 0
ind <- which(haart2[, "death"] == 1)
time <- as.numeric(haart2[ind, "date.death"] - haart2[ind, "init.date"])
temp <- data.frame(ind, time)
haart2[temp[, "ind"], "death.1yr"] <- as.numeric(temp[, "time"] <= 365)

haart2[ind, "followup"] <- temp[, "time"]
ind.2 <- which(haart2[, "death"] == 0)
time.2 <- as.numeric(haart2[ind.2, "last.visit"] - haart2[ind.2, "init.date"])
temp.2 <- data.frame(ind.2, time.2)
haart2[ind.2, "followup"] <- temp.2[, "time.2"]
for (i in 1:nrow(haart2)) {
  if (haart2[i, "followup"] >= 365) {
    haart2[i, "followup"] = 365
  }
}

for (i in 1:nrow(haart2)) {
```

```

    if (haart2[i, "followup"] < 365 & haart2[i, "death"] == 0) {
      haart2[i, "lossfu"] = 1
    } else {
      haart2[i, "lossfu"] = 0
    }
  }
}

```

```
haart2[, all.reg] <- 0
```

```

for (i in 1:nrow(haart2)) {
  reg <- unlist(strsplit(haart2[i, "init.reg"], ","))
  haart2[i, reg] <- 1
}

```

```

new <- rbind(haart, haart2)
head(new)[1:5, ]

```

```

##   male age aids cd4baseline logvl  weight hemoglobin  init.reg
## 1    1  25   0         NA    NA      NA          NA 3TC,AZT,EFV
## 2    1  49   0        143    NA  58.0608         11 3TC,AZT,EFV
## 3    1  42   1        102    NA  48.0816          1 3TC,AZT,EFV
## 4    0  33   0        107    NA  46.0000         NA 3TC,AZT,NVP
## 5    1  27   0         52     4     NA          NA 3TC,D4T,EFV
##   init.date last.visit death date.death death.1yr followup lossfu 3TC AZT
## 1 2003-07-01 2007-02-26     0      <NA>         0     365     0   1   1
## 2 2004-11-23 2008-02-22     0      <NA>         0     365     0   1   1
## 3 2003-04-30 2005-11-21     1 2006-01-11         0     365     0   1   1
## 4 2006-03-25 2006-05-05     1 2006-05-07         1      43     0   1   1
## 5 2004-09-01 2007-11-13     0      <NA>         0     365     0   1   0
##   EFV NVP D4T ABC DDI IDV LPV RTV SQV FTC TDF DDC NFV T20 ATV FPV
## 1    1    0    0    0    0    0    0    0    0    0    0    0    0    0    0
## 2    1    0    0    0    0    0    0    0    0    0    0    0    0    0    0
## 3    1    0    0    0    0    0    0    0    0    0    0    0    0    0    0
## 4    0    1    0    0    0    0    0    0    0    0    0    0    0    0    0
## 5    1    0    1    0    0    0    0    0    0    0    0    0    0    0    0

```

```
tail(new)[2:6, ]
```

```

##      male      age aids cd4baseline  logvl  weight hemoglobin
## 1000    0 40.00000    1        131    NA  46.2672          8
## 1001    0 27.00000    0        232    NA     NA          NA
## 1002    1 38.72142    0        170    NA  84.0000          NA
## 1003    1 23.00000   NA        154 3.995635 65.5000         14
## 1004    0 31.00000    0        236    NA  45.8136          NA
##      init.reg init.date last.visit death date.death death.1yr followup
## 1000 3TC,D4T,NVP 2003-07-03 2008-02-29     0      <NA>         0     365
## 1001 3TC,AZT,NVP 2003-12-01 2004-01-05     0      <NA>         0      35
## 1002 3TC,AZT,NVP 2002-09-26 2004-03-29     0      <NA>         0     365

```

```
## 1003 3TC,DDI,EFV 2007-01-31 2007-04-16 0 <NA> 0 75
## 1004 3TC,D4T,NVP 2003-12-03 2007-10-11 0 <NA> 0 365
##      lossfu 3TC AZT EFV NVP D4T ABC DDI IDV LPV RTV SQV FTC TDF DDC NFV
## 1000      0  1  0  0  1  1  0  0  0  0  0  0  0  0  0  0
## 1001      1  1  1  0  1  0  0  0  0  0  0  0  0  0  0  0
## 1002      0  1  1  0  1  0  0  0  0  0  0  0  0  0  0  0
## 1003      1  1  0  1  0  0  0  1  0  0  0  0  0  0  0  0
## 1004      0  1  0  0  1  1  0  0  0  0  0  0  0  0  0  0
##      T20 ATV FPV
## 1000      0  0  0
## 1001      0  0  0
## 1002      0  0  0
## 1003      0  0  0
## 1004      0  0  0
```

Question 2

14 points

Use the following code to generate data for patients with repeated measures of A1C (a test for levels of blood glucose).

```
genData <- function(n) {
  if (exists(".Random.seed", envir = .GlobalEnv)) {
    save.seed <- get(".Random.seed", envir = .GlobalEnv)
    on.exit(assign(".Random.seed", save.seed, envir = .GlobalEnv))
  } else {
    on.exit(rm(".Random.seed", envir = .GlobalEnv))
  }
  set.seed(n)
  subj <- ceiling(n/10)
  id <- sample(subj, n, replace = TRUE)
  times <- as.integer(difftime(as.POSIXct("2005-01-01"), as.POSIXct("2000-01-01"),
    units = "secs"))
  dt <- as.POSIXct(sample(times, n), origin = "2000-01-01")
  mu <- runif(subj, 4, 10)
  a1c <- unsplit(mapply(rnorm, tabulate(id), mu, SIMPLIFY = FALSE), id)
  data.frame(id, dt, a1c)
}
```

Perform the following manipulations: (2 points each)

1. Order the data set by `id` and `dt`.

```
x <- genData(500)
x <- x[order(x[, "id"], x[, "dt"]), ]
```

2. For each `id`, determine if there is more than a one year gap in
between observations. Add a new row at the one year mark, with the `a1c`
value set to missing. A two year gap would require two new rows, and so
forth.

```

numobs <- nrow(x)

date1 <- x[1, "dt"]
curr.id <- x[1, "id"]

for (i in 2:nrow(x)) {
  if (x[i, "id"] != curr.id) {
    date1 = x[i, "dt"]
    curr.id <- x[i, "id"]
  } else {

    date2 <- x[i, "dt"]
    date2
    timegap <- as.numeric(date2 - date1)
    if (timegap > 365 & timegap <= 730) {
      missing <- as.Date(date1, format = "%Y-%m-%d") + 365
      x[numobs + 1, "dt"] = missing
      x[numobs + 1, "id"] = curr.id
      x[numobs + 1, "a1c"] = "."
      numobs = numobs + 1
    } else if (timegap > 730) {
      missing <- as.Date(date1, format = "%Y-%m-%d") + 365
      x[numobs + 1, "dt"] = missing
      x[numobs + 1, "id"] = curr.id
      x[numobs + 1, "a1c"] = "."
      missing2 <- as.Date(date1, format = "%Y-%m-%d") + 730
      x[numobs + 2, "dt"] = missing2
      x[numobs + 2, "id"] = curr.id
      x[numobs + 2, "a1c"] = "."
      numobs = numobs + 2
    }
    date1 = date2
  }
}

x <- x[order(x[, "id"], x[, "dt"]), ]

```

3. Create a new column `visit`. For each `id`, add the visit number.
 ##### This should be 1 to `n` where `n` is the number of observations for an
 ##### individual. This should include the observations created with missing a1c
 ##### values.

```

curr.id <- x[1, "id"]
n = 1

for (i in 1:nrow(x)) {
  if (x[i, "id"] == curr.id) {
    x[i, "visit"] = n
    n = n + 1
  } else if (x[i, "id"] != curr.id) {
    curr.id <- x[i, "id"]
  }
}

```

```

        x[i, "visit"] = 1
        n = 2
    }
}

num.ind <- curr.id

##### 4. For each `id`, replace missing values with the mean `a1c` value for
##### that individual.

temp <- x

for (i in 1:nrow(x)) {
    if (temp[i, "a1c"] == ".") {
        temp[i, "a1c"] = 0
    }
}

curr.id <- temp[1, "id"]
mean <- numeric(num.ind)
total.visits <- numeric(num.ind)
mean.t <- as.numeric(temp[1, "a1c"])
num.miss <- 0

for (i in 2:nrow(temp)) {
    if (temp[i, "id"] == curr.id) {
        mean.t = mean.t + as.numeric(temp[i, "a1c"])
        if (as.numeric(temp[i, "a1c"]) == 0) {
            num.miss = num.miss + 1
        }
    } else if (temp[i, "id"] != curr.id) {
        mean[curr.id] <- mean.t/(temp[i - 1, "visit"] - num.miss)
        total.visits[curr.id] <- temp[i - 1, "visit"]
        curr.id <- temp[i, "id"]
        num.miss <- 0
        mean.t <- as.numeric(temp[i, "a1c"])
    }
    if (i == nrow(temp)) {
        mean[curr.id] <- mean.t/(temp[i, "visit"] - num.miss)
        total.visits[curr.id] <- temp[i, "visit"]
    }
}
mean

```

```

## [1] 4.063372 7.544643 6.757640 3.892127 9.512311 7.555965 9.161686
## [8] 7.189064 9.283873 7.975217 6.917562 7.034021 9.145282 6.623756
## [15] 8.012406 4.222158 3.996034 9.164873 5.507210 3.726675 8.140939
## [22] 5.637501 7.366889 7.439316 6.877135 6.556759 4.926457 7.433917
## [29] 4.508086 6.045577 7.116586 6.568791 6.494069 6.768615 8.476700
## [36] 9.604410 9.606253 5.355979 6.917013 9.530136 9.802424 3.891770
## [43] 6.095849 9.091670 6.737204 9.621763 9.231489 6.404600 6.096076
## [50] 8.962319

```



```

for (i in 1:nrow(x)) {
  if (x[i, "a1c"] == ".") {
    x[i, "a1c"] = mean[x[i, "id"]]
  }
}

##### 5. Print mean `a1c` for each `id`. & ##### 6. Print total number of
##### visits for each `id`.

(y <- data.frame(mean, total.visits))

```

```

##          mean total.visits
## 1  4.063372           11
## 2  7.544643           20
## 3  6.757640           14
## 4  3.892127           12
## 5  9.512311           14
## 6  7.555965           10
## 7  9.161686            9
## 8  7.189064           12
## 9  9.283873           11
## 10 7.975217           12
## 11 6.917562           10
## 12 7.034021           10
## 13 9.145282            8
## 14 6.623756           12
## 15 8.012406            8
## 16 4.222158            9
## 17 3.996034           12
## 18 9.164873           10
## 19 5.507210           10
## 20 3.726675            9
## 21 8.140939           10
## 22 5.637501            8
## 23 7.366889            8
## 24 7.439316           15
## 25 6.877135           12
## 26 6.556759           14
## 27 4.926457           11
## 28 7.433917           14
## 29 4.508086           10
## 30 6.045577            7
## 31 7.116586           11
## 32 6.568791            5
## 33 6.494069            8
## 34 6.768615           12
## 35 8.476700           11
## 36 9.604410            9
## 37 9.606253           17
## 38 5.355979           15
## 39 6.917013            8
## 40 9.530136            7

```

```
## 41 9.802424      17
## 42 3.891770      14
## 43 6.095849      11
## 44 9.091670      11
## 45 6.737204      14
## 46 9.621763       9
## 47 9.231489      12
## 48 6.404600      11
## 49 6.096076      12
## 50 8.962319      10
```

```
##### 7. Print the observations for `id = 15`.
```

```
x[which(x[, "id"] == 15), ]
```

```
##      id      dt      a1c visit
## 11  15 2000-04-30 00:34:50 7.52710515747364      1
## 406 15 2001-01-17 21:11:02 5.89837126480442      2
## 306 15 2001-04-25 06:23:05 8.56659306505127      3
## 518 15 2002-04-24 19:00:00 8.01240569465381      4
## 519 15 2003-04-24 19:00:00 8.01240569465381      5
## 484 15 2003-06-06 14:06:00 9.13376871828962      6
## 520 15 2004-06-04 19:00:00 8.01240569465381      7
## 263 15 2004-08-20 17:47:11 8.93619026765011      8
```

Question 3

10 points

Import the `addr.txt` file from the GitHub repository. This file contains a listing of names and addresses (thanks google). Parse each line to create a data.frame with the following columns: lastname, firstname, streetno, streetname, city, state, zip. Keep middle initials or abbreviated names in the firstname column. Print out the entire data.frame.

```
addr <- read.csv("https://raw.githubusercontent.com/fonnesbeck/Bios6301/master/datasets/addr.txt",
  stringsAsFactors = FALSE)
col <- c("lastname", "firstname", "streetno", "streetname", "city", "state",
  "zip")
for (i in 1:length(col)) {
  addr[, col[i]] <- character(nrow(addr))
}

for (i in 1:nrow(addr)) {
  x.temp <- unlist(strsplit(addr[i, 1], " "))
  x <- x.temp[x.temp != ""]
  addr[i, "lastname"] = x[1]
  addr[i, "firstname"] = x[2]
  addr[i, "streetno"] = gsub("([0-9]{1,4}).*", "\\1", x[3])
  addr[i, "streetname"] = gsub("[0-9]{1,4} (.*)", "\\1", x[3])
  addr[i, "city"] = x[4]
  addr[i, "state"] = x[5]
  addr[i, "zip"] = x[6]
}
```

```
addr[, "zip"] = gsub("0", "0", addr[, "zip"])
addr[, 1] = NULL
```

```
addr
```

##	lastname	firstname	streetno	streetname	city	state
## 1	Barnaby	David	373	W. Geneva St.	Wms. Bay	WI
## 2	Bausch	Judy	373	W. Geneva St.	Wms. Bay	WI
## 3	Bolatto	Alberto	725	Commonwealth Ave.	Boston	MA
## 4	Carlstrom	John	933	E. 56th St.	Chicago	IL
## 5	Chamberlin	Richard A.	111	Nowelo St.	Hilo	HI
## 6	Chuss	Dave	2145	Sheridan Rd	Evanston	IL
## 7	Davis	E. J.	933	E. 56th St.	Chicago	IL
## 8	Depoy	Darren	174	W. 18th Ave.	Columbus	OH
## 9	Griffin	Greg	5000	Forbes Ave.	Pittsburgh	PA
## 10	Halvorsen	Nils	933	E. 56th St.	Chicago	IL
## 11	Harper	Al	373	W. Geneva St.	Wms. Bay	WI
## 12	Huang	Maohai	725	W. Commonwealth Ave.	Boston	MA
## 13	Ingalls	James G.	725	W. Commonwealth Ave.	Boston	MA
## 14	Jackson	James M.	725	W. Commonwealth Ave.	Boston	MA
## 15	Knudsen	Scott	373	W. Geneva St.	Wms. Bay	WI
## 16	Kovac	John	5640	S. Ellis Ave.	Chicago	IL
## 17	Landsberg	Randy	5640	S. Ellis Ave.	Chicago	IL
## 18	Lo	Kwok-Yung	1002	W. Green St.	Urbana	IL
## 19	Loewenstein	Robert F.	373	W. Geneva St.	Wms. Bay	WI
## 20	Lynch	John	4201	Wilson Blvd	Arlington	VA
## 21	Martini	Paul	174	W. 18th Ave.	Columbus	OH
## 22	Meyer	Stephan	933	E. 56th St.	Chicago	IL
## 23	Mrozek	Fred	373	W. Geneva St.	Wms. Bay	WI
## 24	Newcomb	Matt	5000	Forbes Ave.	Pittsburgh	PA
## 25	Novak	Giles	2145	Sheridan Rd	Evanston	IL
## 26	Odalen	Nancy	373	W. Geneva St.	Wms. Bay	WI
## 27	Pernic	Dave	373	W. Geneva St.	Wms. Bay	WI
## 28	Pernic	Bob	373	W. Geneva St.	Wms. Bay	WI
## 29	Peterson	Jeffrey	5000	Forbes Ave.	Pittsburgh	PA
## 30	Pryke	Clem	933	E. 56th St.	Chicago	IL
## 31	Rebull	Luisa	5640	S. Ellis Ave.	Chicago	IL
## 32	Renbarger	Thomas	2145	Sheridan Rd	Evanston	IL
## 33	Rottman	Joe	8730	W. Mountain View Ln	Littleton	CO
## 34	Schartman	Ethan	933	E. 56th St.	Chicago	IL
## 35	Spotz	Bob	373	W. Geneva St.	Wms. Bay	WI
## 36	Thoma	Mark	373	W. Geneva St.	Wms. Bay	WI
## 37	Walker	Chris	933	N. Cherry St.	Tucson	AZ
## 38	Wehrer	Cheryl	5000	Forbes Ave.	Pittsburgh	PA
## 39	Wirth	Jesse	373	W. Geneva St.	Wms. Bay	WI
## 40	Wright	Greg	791	Holmdel-Keyport Rd.	Holmdel	NY
## 41	Zingale	Michael	5640	S. Ellis Ave.	Chicago	IL
##	zip					
## 1	53191					
## 2	53191					
## 3	02215					
## 4	60637					
## 5	96720					

```
## 6 60208-3112
## 7 60637
## 8 43210
## 9 15213
## 10 60637
## 11 53191
## 12 02215
## 13 02215
## 14 02215
## 15 53191
## 16 60637
## 17 60637
## 18 61801
## 19 53191
## 20 22230
## 21 43210
## 22 60637
## 23 53191
## 24 15213
## 25 60208-3112
## 26 53191
## 27 53191
## 28 53191
## 29 15213
## 30 60637
## 31 60637
## 32 60208-3112
## 33 80125
## 34 60637
## 35 53191
## 36 53191
## 37 85721
## 38 15213
## 39 53191
## 40 07733-1988
## 41 60637
```

Question 4

2 points

The first argument to most functions that fit linear models are formulas. The following example defines the response variable `death` and allows the model to incorporate all other variables as terms. `.` is used to mean all columns not otherwise in the formula.

```
url <- "https://github.com/fonnesbeck/Bios6301/raw/master/datasets/haart.csv"
haart_df <- read.csv(url)[, c("death", "weight", "hemoglobin", "cd4baseline")]
coef(summary(glm(death ~ ., data = haart_df, family = binomial(logit))))
```

```
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  3.576411744 1.226870535  2.915069 0.0035561039
## weight      -0.046210552 0.022556001 -2.048703 0.0404911395
## hemoglobin   -0.350642786 0.105064078 -3.337418 0.0008456055
## cd4baseline  0.002092582 0.001811959  1.154872 0.2481427160
```

Now imagine running the above several times, but with a different response and data set each time. Here's a function:

```
myfun <- function(dat, response) {  
  form <- as.formula(response ~ .)  
  coef(summary(glm(form, data = dat, family = binomial(logit))))  
}
```

Unfortunately, it doesn't work. `tryCatch` is "catching" the error so that this file can be knit to PDF.

```
tryCatch(myfun(haart_df, death), error = function(e) e)
```

```
## <simpleError in eval(expr, envir, enclos): object 'death' not found>
```

What do you think is going on? Consider using `debug` to trace the problem.

```
debug(myfun)  
myfun(haart_df, death)  
undebug(myfun)
```

The error is: 'object 'death' not found'

I believe that the issue is that we are trying to pass into the function something which is to be used as a string, but the function is expecting it to be a variable.

5 bonus points

Create a working function.

```
myfun3 <- function(dat, response) {  
  form <- paste("as.formula(", response, "~ .)")  
  coef(summary(glm(form, data = dat, family = binomial(logit))))  
}  
myfun3(haart_df, "death")
```

```
##               Estimate Std. Error z value Pr(>|z|)  
## (Intercept)  3.576411744 1.226870535  2.915069 0.0035561039  
## weight      -0.046210552 0.022556001 -2.048703 0.0404911395  
## hemoglobin  -0.350642786 0.105064078 -3.337418 0.0008456055  
## cd4baseline  0.002092582 0.001811959  1.154872 0.2481427160
```