Getting Started with R

R Handout

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Always First Command

```
> library(NCStats)
```

Basic Data Frames

```
> setwd("C:/aaaWork/Web/GitHub/NCMTH107/modules/Getting_Started_R")
> iris <- read.csv("Iris.csv")
> str(iris)

'data.frame': 150 obs. of 5 variables:
$ seplen : int 50 46 46 51 55 48 52 49 44 50 ...
$ sepwid : int 33 34 36 33 35 31 34 36 32 35 ...
$ petlen : int 14 14 10 17 13 16 14 14 13 16 ...
$ petwid : int 2 3 2 5 2 2 2 1 2 6 ...
$ species: Factor w/ 3 levels "setosa","versicolor",..: 1 1 1 1 1 1 1 1 1 1 ...
> headtail(iris) # NOT the entire data.frame
```

```
seplen sepwid petlen petwid
                             species
1
       50
             33
                   14
                           2
                               setosa
2
       46
             34
                   14
                           3
                               setosa
3
       46
             36
                   10
                          2
                               setosa
148
       57
             25
                   50
                          20 virginica
             28
149
       58
                   51
                          24 virginica
150
       63
             33
                   60
                          25 virginica
```

```
> ls()
```

```
[1] "f" "fnm" "gt5" "iris" "need2render" "set" [7] "setver" "ver_and_gt5" "ver_or_gt5"
```

Working With Data Frames

```
> iris[2,]
```

```
seplen sepwid petlen petwid species
      46
             34
                    14
                            3 setosa
> iris$seplen
  [1] 50 46 46 51 55 48 52 49 44 50 44 47 48 51 48 50 50 43 58 51 49 51 50 46 57 50 54 52
 [29] 55 49 54 50 44 47 46 51 50 49 54 54 51 48 48 45 57 51 54 51 52 53 65 62 59 61 60 56
 [57] 57 63 70 64 61 55 54 58 55 50 67 56 58 60 57 57 49 56 57 66 52 60 50 55 58 62 59 60
 [85] 67 63 56 63 61 64 51 57 61 56 69 55 55 66 68 67 64 67 63 69 65 65 58 68 62 77 67 76
[113] 49 67 59 63 64 79 67 77 63 72 61 61 64 62 77 63 58 72 71 64 60 63 77 60 69 74 56 73
[141] 67 65 69 72 65 64 68 57 58 63
> iris$seplen[3]
[1] 46
> iris[3,2]
[1] 36
> iris[c(1,3,5),]
  seplen sepwid petlen petwid species
             33
                    14
                            2 setosa
3
      46
             36
                    10
                            2 setosa
5
      55
             35
                    13
                            2 setosa
Filtering Data Frames
> set <- filterD(iris,species=="setosa")</pre>
> str(set)
'data.frame':
              50 obs. of 5 variables:
 $ seplen : int 50 46 46 51 55 48 52 49 44 50 ...
 $ sepwid : int 33 34 36 33 35 31 34 36 32 35 ...
 $ petlen : int 14 14 10 17 13 16 14 14 13 16 ...
 $ petwid : int  2 3 2 5 2 2 2 1 2 6 ...
 $ species: Factor w/ 1 level "setosa": 1 1 1 1 1 1 1 1 1 1 ...
> gt5 <- filterD(iris,seplen>5)
> headtail(gt5)
    seplen sepwid petlen petwid
                                  species
               33
                              2
1
        50
                      14
                                   setosa
2
        46
               34
                      14
                              3
                                   setosa
               36
                      10
                             2
3
        46
                                   setosa
148
       57
               25
                      50
                             20 virginica
```

24 virginica

25 virginica

149

150

58

63

28

33

51

60

```
> setver1 <- filterD(iris,species!="virginica")</pre>
> str(setver1)
                100 obs. of 5 variables:
'data.frame':
 $ seplen : int 50 46 46 51 55 48 52 49 44 50 ...
 $ sepwid : int 33 34 36 33 35 31 34 36 32 35 ...
 $ petlen : int 14 14 10 17 13 16 14 14 13 16 ...
 \ petwid : int \ 2 3 2 5 2 2 2 1 2 6 ...
 $ species: Factor w/ 2 levels "setosa", "versicolor": 1 1 1 1 1 1 1 1 1 1 ...
> setver <- filterD(iris, species %in% c("setosa", "versicolor"))</pre>
> str(setver)
'data.frame':
              100 obs. of 5 variables:
$ seplen : int 50 46 46 51 55 48 52 49 44 50 ...
\ sepwid : int \ 33 34 36 33 35 31 34 36 32 35 ...
 $ petlen : int 14 14 10 17 13 16 14 14 13 16 ...
 $ petwid : int  2 3 2 5 2 2 2 1 2 6 ...
 $ species: Factor w/ 2 levels "setosa", "versicolor": 1 1 1 1 1 1 1 1 1 1 ...
> ver_and_gt5 <- filterD(iris,species=="versicolor",seplen>5)
> headtail(ver and gt5)
   seplen sepwid petlen petwid
                                  species
1
              28
                    46
                            15 versicolor
2
      62
              22
                     45
                            15 versicolor
3
      59
              32
                     48
                            18 versicolor
48
      66
              30
                     44
                           14 versicolor
49
      68
              28
                     48
                           14 versicolor
50
      67
              30
                     50
                           17 versicolor
> ver_or_gt5 <- filterD(iris,species=="versicolor" | seplen>5)
> headtail(ver_or_gt5)
    seplen sepwid petlen petwid
                                  species
               33
1
        50
                      14
                                   setosa
2
        46
               34
                      14
                                   setosa
                              3
               36
3
        46
                      10
                             2
                                   setosa
        57
               25
                      50
148
                             20 virginica
149
        58
               28
                      51
                             24 virginica
150
               33
                             25 virginica
       63
                      60
```