

Sta-518 Self Reflection

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STA 518 Objectives:-

1.Import, manage, and clean data:-

I can import data from a variety of sources.

```
# Read TXT files with read.table()
children <- read.table("https://alexdl06.github.io/intro2R/data/children.txt", header = TRUE)
children
```

```
##      names sex age weight height
## 1  ALFRED  M  14     69    112
## 2 BARBARA  F  13     62    102
## 3   JAMES  M  12     57     83
## 4   JANE   F  12     59     84
## 5   JOHN   M  12     59     99
## 6   JUDY   F  14     64     90
## 7 LOUISE   F  12     56     77
## 8   MARY   F  15     66    112
## 9  RONALD  M  15     67    133
## 10 WILLIAM M  15     66    112
```

```
# Read in csv files with read_csv()
college<- read_csv("data/recent-grads.csv")
```

```
##
## -- Column specification -----
## cols(
##   .default = col_double(),
##   major = col_character(),
##   major_category = col_character()
## )
## i Use `spec()` for the full column specifications.
college
```

```
## # A tibble: 173 x 21
##   rank major_code major          major_category total sample_size  men women
##   <dbl>    <dbl> <chr>          <chr>          <dbl>    <dbl> <dbl> <dbl>
## 1     1        2419 Petroleum Engi~ Engineering    2339        36  2057   282
## 2     2        2416 Mining And Min~ Engineering     756         7   679    77
## 3     3        2415 Metallurgical ~ Engineering     856         3   725   131
## 4     4        2417 Naval Architec~ Engineering    1258        16  1123   135
## 5     5        2405 Chemical Engin~ Engineering   32260       289 21239 11021
## 6     6        2418 Nuclear Engine~ Engineering    2573        17  2200   373
```

```
## 7      7      6202 Actuarial Scie~ Business      3777      51 2110 1667
## 8      8      5001 Astronomy And ~ Physical Scie~ 1792      10 832 960
## 9      9      2414 Mechanical Eng~ Engineering 91227      1029 80320 10907
## 10     10      2408 Electrical Eng~ Engineering 81527      631 65511 16016
## # ... with 163 more rows, and 13 more variables: sharewomen <dbl>,
## #   employed <dbl>, employed_fulltime <dbl>, employed_parttime <dbl>,
## #   employed_fulltime_yearround <dbl>, unemployed <dbl>,
## #   unemployment_rate <dbl>, p25th <dbl>, median <dbl>, p75th <dbl>,
## #   college_jobs <dbl>, non_college_jobs <dbl>, low_wage_jobs <dbl>
```

```
# Read in xlsx files with read_excel()
fxlsx <- read_excel("data/loyn.xlsx")
fxlsx
```

```
## # A tibble: 56 x 8
##   Site ABUND AREA DIST LDIST YR.ISOL GRAZE ALT
##   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1     1   5.3  0.1   39    39   1968     2   160
## 2     2     2  0.5  234   234   1920     5    60
## 3     3   1.5  0.5  104   311   1900     5   140
## 4     4  17.1  1     66    66   1966     3   160
## 5     5  13.8  1    246   246   1918     5   140
## 6     6  14.1  1    234   285   1965     3   130
## 7     7   3.8  1    467   467   1955     5    90
## 8     8   2.2  1    284  1829   1920     5    60
## 9     9   3.3  1    156   156   1965     4   130
## 10    10     3  1    311   571   1900     5   130
## # ... with 46 more rows
```

```
# Read in csv files with readr::read_csv()
banksal <- readr::read_csv("data/banksalary.csv")
```

```
##
## -- Column specification -----
## cols(
##   bsal = col_double(),
##   ansal = col_double(),
##   sex = col_character(),
##   senior = col_double(),
##   age = col_double(),
##   educ = col_double(),
##   exper = col_double()
## )
```

```
banksal
```

```
## # A tibble: 93 x 7
##   bsal ansal sex senior age educ exper
##   <dbl> <dbl> <chr> <dbl> <dbl> <dbl> <dbl>
## 1 5040 12420 MALE     96 329 15 14
## 2 6300 12060 MALE     82 357 15 72
## 3 6000 15120 MALE     67 315 15 35.5
## 4 6000 16320. MALE     97 354 12 24
## 5 6000 12300 MALE     66 351 12 56
## 6 6840 10380 MALE     92 374 15 41.5
## 7 8100 13980. MALE     66 369 16 54.5
## 8 6000 10140 MALE     82 363 12 32
```

```
## 9 6000 12360 MALE      88 555 12 252
## 10 6900 10920 MALE      75 416 15 132
## # ... with 83 more rows
```

```
# reading a large data set
```

```
nls <- read_dta(file="https://github.com/ozanj/rclass/raw/master/data/nls72/nls72stu_percontor_vars.dta")
nls
```

```
## # A tibble: 22,652 x 89
```

```
##       id schcode  bysex  srfq2a  srfq2d  fsex schlmrkr  ssex  crace
##   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1    18   3000 2 [2. f~ 98 [98. ~ 98 [98. ~ NA      1 [1. 1~ 1 [1. ~ 4 [4. ~
## 2    67   3000 1 [1. m~ 98 [98. ~ 98 [98. ~ NA      0 [0. N~ 1 [1. ~ 2 [2. ~
## 3    83   2518 2 [2. f~ 71 [71. ~ 98 [98. ~ 2 [2. ~ 1 [1. 1~ 99 [99.~ 7 [7. ~
## 4   174   2911 1 [1. m~ 98 [98. ~ 71 [71. ~ NA      1 [1. 1~ 1 [1. ~ 7 [7. ~
## 5   190    800 2 [2. f~ 98 [98. ~ 98 [98. ~ 2 [2. ~ 1 [1. 1~ 2 [2. ~ 7 [7. ~
## 6   232   7507 2 [2. f~ 98 [98. ~ 98 [98. ~ 2 [2. ~ 1 [1. 1~ 2 [2. ~ 7 [7. ~
## 7   315   3000 1 [1. m~ 72 [72. ~ 71 [71. ~ NA      0 [0. N~ 1 [1. ~ 7 [7. ~
## 8   380   9516 1 [1. m~ 98 [98. ~ 98 [98. ~ NA      1 [1. 1~ 1 [1. ~ 7 [7. ~
## 9   414   2518 1 [1. m~ 98 [98. ~ 98 [98. ~ NA      0 [0. N~ 98 [98.~ 7 [7. ~
## 10  430   2701 1 [1. m~ 98 [98. ~ 98 [98. ~ NA      1 [1. 1~ 1 [1. ~ 4 [4. ~
## # ... with 22,642 more rows, and 80 more variables: csex <dbl>,
## #   tfusex <dbl>, ft67 <dbl>, ftfusex <dbl>, fi29 <dbl>,
## #   fi39 <dbl>, fi43 <dbl>, fi52a <dbl>, fi52b <dbl>,
## #   fi52c <dbl>, fi52d <dbl>, fi52e <dbl>, fi52f <dbl>,
## #   race86 <dbl>, birthmon <dbl>, birthday <dbl>, birthyr <dbl>,
## #   srfq0 <dbl>, srfq2b <dbl>, srfq2c <dbl>, srfq2e <dbl>, srfq2f <dbl>,
## #   srfq2g <dbl>, srfq2h <dbl>, srfq2i <dbl>, scvocsc <dbl>, scpic1 <dbl>,
## #   scpic2 <dbl>, scpict <dbl>, scrdsc <dbl>, sclgsc <dbl>, scmatsc <dbl>,
## #   scm scm1 <dbl>, scm scm2 <dbl>, scm scm3 <dbl>, scm scm4 <dbl>, srifprk <dbl>,
## #   fbirthmo <dbl>, fbirthda <dbl>, fbirthyr <dbl>, cfaocp <dbl>,
## #   cbirthm <dbl>, cbirthd <dbl>, cbirthyr <dbl>, fi40cm <dbl>, fi40cy <dbl>,
## #   fi41cm <dbl>, fi41cy <dbl>, fi42cm <dbl>, fi42by <dbl>, fi111a84 <dbl>,
## #   fi111b84 <dbl>, fi111c84 <dbl>, fi111d84 <dbl>, fi111e84 <dbl>,
## #   fi111f84 <dbl>, fi111g84 <dbl>, fi111h84 <dbl>, fi111i84 <dbl>,
## #   fi111j84 <dbl>, fi111k84 <dbl>, fi111l84 <dbl>, fi111m84 <dbl>,
## #   fi111n84 <dbl>, fi111o84 <dbl>, fi111a85 <dbl>, fi111b85 <dbl>,
## #   fi111c85 <dbl>, fi111d85 <dbl>, fi111e85 <dbl>, fi111f85 <dbl>,
## #   fi111g85 <dbl>, fi111h85 <dbl>, fi111i85 <dbl>, fi111j85 <dbl>,
## #   fi111k85 <dbl>, fi111l85 <dbl>, fi111m85 <dbl>, fi111n85 <dbl>,
## #   fi111o85 <dbl>
```

```
nc <- read_csv("data/nc.csv")
```

```
##
## -- Column specification -----
## cols(
##   fage = col_double(),
##   mage = col_double(),
##   mature = col_character(),
##   weeks = col_double(),
##   premie = col_character(),
##   visits = col_double(),
##   marital = col_character(),
##   gained = col_double(),
##   weight = col_double(),
```

```
## lowbirthweight = col_character(),
## gender = col_character(),
## habit = col_character(),
## whitemom = col_character()
## )

country <- read_excel("data/country.xlsx")
people <- read_excel("data/people.xlsx")
```

I can isolate information from a larger data source.

```
# I am taking the above children table to perform data isolation operations
# filter operation
```

```
filter(children, sex == "M" & age == "15")
```

```
##      names sex age weight height
## 1  RONALD   M  15      67     133
## 2 WILLIAM   M  15      66     112
```

```
# I am taking the above banksal table to perform data isolation operations
```

```
fbanksal <- filter(banksal, bsal >= 6000 & senior >= 60)
fbanksal
```

```
## # A tibble: 26 x 7
##   bsal  ansal sex  senior  age  educ exper
##   <dbl> <dbl> <chr>   <dbl> <dbl> <dbl> <dbl>
## 1  6300 12060 MALE      82   357    15    72
## 2  6000 15120 MALE      67   315    15   35.5
## 3  6000 16320 MALE      97   354    12    24
## 4  6000 12300 MALE      66   351    12    56
## 5  6840 10380 MALE      92   374    15   41.5
## 6  8100 13980 MALE      66   369    16   54.5
## 7  6000 10140 MALE      82   363    12    32
## 8  6000 12360 MALE      88   555    12   252
## 9  6900 10920 MALE      75   416    15   132
## 10 6900 10920 MALE      89   481    12   175
## # ... with 16 more rows
```

```
# Randomly select rows from banksal table
```

```
slice_sample(banksal, n = 5, replace = TRUE)
```

```
## # A tibble: 5 x 7
##   bsal  ansal sex  senior  age  educ exper
##   <dbl> <dbl> <chr>   <dbl> <dbl> <dbl> <dbl>
## 1  5040 12420 MALE      96   329    15    14
## 2  6300  9780 FEMALE    66   394    12   86.5
## 3  8100 13980 MALE      66   369    16   54.5
## 4  6300 10860 FEMALE    84   662    15   231
## 5  5400 12600 MALE      78   305    12     7
```

```
# using select statement
```

```
select(iris, petal_length = Petal.Length)
```

```
##      petal_length
## 1              1.4
## 2              1.4
## 3              1.3
```

## 4	1.5
## 5	1.4
## 6	1.7
## 7	1.4
## 8	1.5
## 9	1.4
## 10	1.5
## 11	1.5
## 12	1.6
## 13	1.4
## 14	1.1
## 15	1.2
## 16	1.5
## 17	1.3
## 18	1.4
## 19	1.7
## 20	1.5
## 21	1.7
## 22	1.5
## 23	1.0
## 24	1.7
## 25	1.9
## 26	1.6
## 27	1.6
## 28	1.5
## 29	1.4
## 30	1.6
## 31	1.6
## 32	1.5
## 33	1.5
## 34	1.4
## 35	1.5
## 36	1.2
## 37	1.3
## 38	1.4
## 39	1.3
## 40	1.5
## 41	1.3
## 42	1.3
## 43	1.3
## 44	1.6
## 45	1.9
## 46	1.4
## 47	1.6
## 48	1.4
## 49	1.5
## 50	1.4
## 51	4.7
## 52	4.5
## 53	4.9
## 54	4.0
## 55	4.6
## 56	4.5
## 57	4.7

## 58	3.3
## 59	4.6
## 60	3.9
## 61	3.5
## 62	4.2
## 63	4.0
## 64	4.7
## 65	3.6
## 66	4.4
## 67	4.5
## 68	4.1
## 69	4.5
## 70	3.9
## 71	4.8
## 72	4.0
## 73	4.9
## 74	4.7
## 75	4.3
## 76	4.4
## 77	4.8
## 78	5.0
## 79	4.5
## 80	3.5
## 81	3.8
## 82	3.7
## 83	3.9
## 84	5.1
## 85	4.5
## 86	4.5
## 87	4.7
## 88	4.4
## 89	4.1
## 90	4.0
## 91	4.4
## 92	4.6
## 93	4.0
## 94	3.3
## 95	4.2
## 96	4.2
## 97	4.2
## 98	4.3
## 99	3.0
## 100	4.1
## 101	6.0
## 102	5.1
## 103	5.9
## 104	5.6
## 105	5.8
## 106	6.6
## 107	4.5
## 108	6.3
## 109	5.8
## 110	6.1
## 111	5.1

```
## 112      5.3
## 113      5.5
## 114      5.0
## 115      5.1
## 116      5.3
## 117      5.5
## 118      6.7
## 119      6.9
## 120      5.0
## 121      5.7
## 122      4.9
## 123      6.7
## 124      4.9
## 125      5.7
## 126      6.0
## 127      4.8
## 128      4.9
## 129      5.6
## 130      5.8
## 131      6.1
## 132      6.4
## 133      5.6
## 134      5.1
## 135      5.6
## 136      6.1
## 137      5.6
## 138      5.5
## 139      4.8
## 140      5.4
## 141      5.6
## 142      5.1
## 143      5.1
## 144      5.9
## 145      5.7
## 146      5.2
## 147      5.0
## 148      5.2
## 149      5.4
## 150      5.1
```

I can combine information from multiple data sources

Example data tables

```
orders <- read.csv("https://raw.githubusercontent.com/ds4stats/r-tutorials/master/merging/data/orders.csv")
orders
```

```
##   order id   date
## 1     1   4 Jan-01
## 2     2   8 Feb-01
## 3     3  42 Apr-15
## 4     4  50 Apr-17
```

```
customers <- read.csv("https://raw.githubusercontent.com/ds4stats/r-tutorials/master/merging/data/customers.csv")
customers
```

```
##   id   name
```

```
## 1 4 Tukey
## 2 8 Wickham
## 3 15 Mason
## 4 16 Jordan
## 5 23 Patil
## 6 42 Cox
```

Joining data tables

Inner_join creates a new table which is restricted to cases where the values of "by variable" exist in both tables.

```
inner_join(x = orders, y = customers, by = "id")
```

```
## order id date name
## 1 1 4 Jan-01 Tukey
## 2 2 8 Feb-01 Wickham
## 3 3 42 Apr-15 Cox
```

Left_join returns all cases from the x data table, regardless of whether there are matching values of the by variable in the y data table.

```
left_join(x = orders, y = customers, by = "id")
```

```
## order id date name
## 1 1 4 Jan-01 Tukey
## 2 2 8 Feb-01 Wickham
## 3 3 42 Apr-15 Cox
## 4 4 50 Apr-17 <NA>
```

Right_join returns all cases from the y data table, regardless of whether there are matching values of the by variable in the x data table.

```
right_join(x = orders, y = customers, by = "id")
```

```
## order id date name
## 1 1 4 Jan-01 Tukey
## 2 2 8 Feb-01 Wickham
## 3 3 42 Apr-15 Cox
## 4 NA 15 <NA> Mason
## 5 NA 16 <NA> Jordan
## 6 NA 23 <NA> Patil
```

Full_join returns all rows and columns from both x and y.

```
full_join(x = orders, y = customers, by = "id")
```

```
## order id date name
## 1 1 4 Jan-01 Tukey
## 2 2 8 Feb-01 Wickham
## 3 3 42 Apr-15 Cox
## 4 4 50 Apr-17 <NA>
## 5 NA 15 <NA> Mason
## 6 NA 16 <NA> Jordan
## 7 NA 23 <NA> Patil
```

Semi_join returns all rows from the x data table where there are matching values of the by variable in the y data table.

```
semi_join(x = orders, y = customers, by = "id")
```

```
## order id date
## 1 1 4 Jan-01
## 2 2 8 Feb-01
## 3 3 42 Apr-15
```

I can restructure information to be in a “tidy” format.


```
# restructure a dataset to be in a more efficient format and add features to make a table more understandable
wide_measures <- country %>%
  filter(year == 2002) %>%
  group_by(continent) %>%
  summarise(
    med_LE = median(pop),
    mean_LE = mean(pop)
  )

wide_measures %>%
  pivot_longer(
    cols = ends_with("_LE"),
    names_to = "measure",
    values_to = "values"
  )
```

```
## # A tibble: 10 x 3
##   continent measure      values
##   <chr>      <chr>      <dbl>
## 1 Africa    med_LE      8821778.
## 2 Africa    mean_LE     16033152.
## 3 Americas  med_LE      8650322
## 4 Americas  mean_LE     33990910.
## 5 Asia      med_LE      22662365
## 6 Asia      mean_LE     109145521.
## 7 Europe    med_LE      9518744
## 8 Europe    mean_LE     19274129.
## 9 Oceania   med_LE      11727414.
## 10 Oceania  mean_LE     11727414.
```

```
long_measures <- wide_measures %>%
  pivot_longer(
    cols = ends_with("_LE"),
    names_to = "measure",
    values_to = "values"
  )

long_measures %>%
  pivot_wider(
    names_from = continent,
    values_from = values
  )
```

```
## # A tibble: 2 x 6
##   measure Africa Americas Asia Europe Oceania
##   <chr>      <dbl>    <dbl>  <dbl>  <dbl>  <dbl>
## 1 med_LE  8821778.  8650322 22662365 9518744 11727414.
## 2 mean_LE 16033152. 33990910. 109145521. 19274129. 11727414.
```

I can transform information to be in a format better suited for specific tasks.

```
# reducing the dataset for better fit
reducing_dataset <- people %>%
  select(skin_color) %>%
  separate(skin_color,
    into = c("skin_first", "skin_second")) %>%
```

```

pivot_longer(cols = everything(),
              names_to = "skin_order",
              values_to = "skin_color",
              values_drop_na = TRUE) %>%
select(skin_color)

```

```
## Warning: Expected 2 pieces. Additional pieces discarded in 4 rows [16, 47, 67,
## 76].
```

```
## Warning: Expected 2 pieces. Missing pieces filled with `NA` in 71 rows [1, 2, 4,
## 5, 6, 7, 9, 10, 11, 12, 13, 14, 15, 17, 18, 19, 20, 21, 22, 23, ...].
```

```
reducing_dataset
```

```

## # A tibble: 103 x 1
##   skin_color
##   <chr>
## 1 fair
## 2 gold
## 3 white
## 4 blue
## 5 white
## 6 light
## 7 light
## 8 light
## 9 white
## 10 red
## # ... with 93 more rows

```

Create graphical displays and numerical summaries of data for exploratory analysis and presentations.

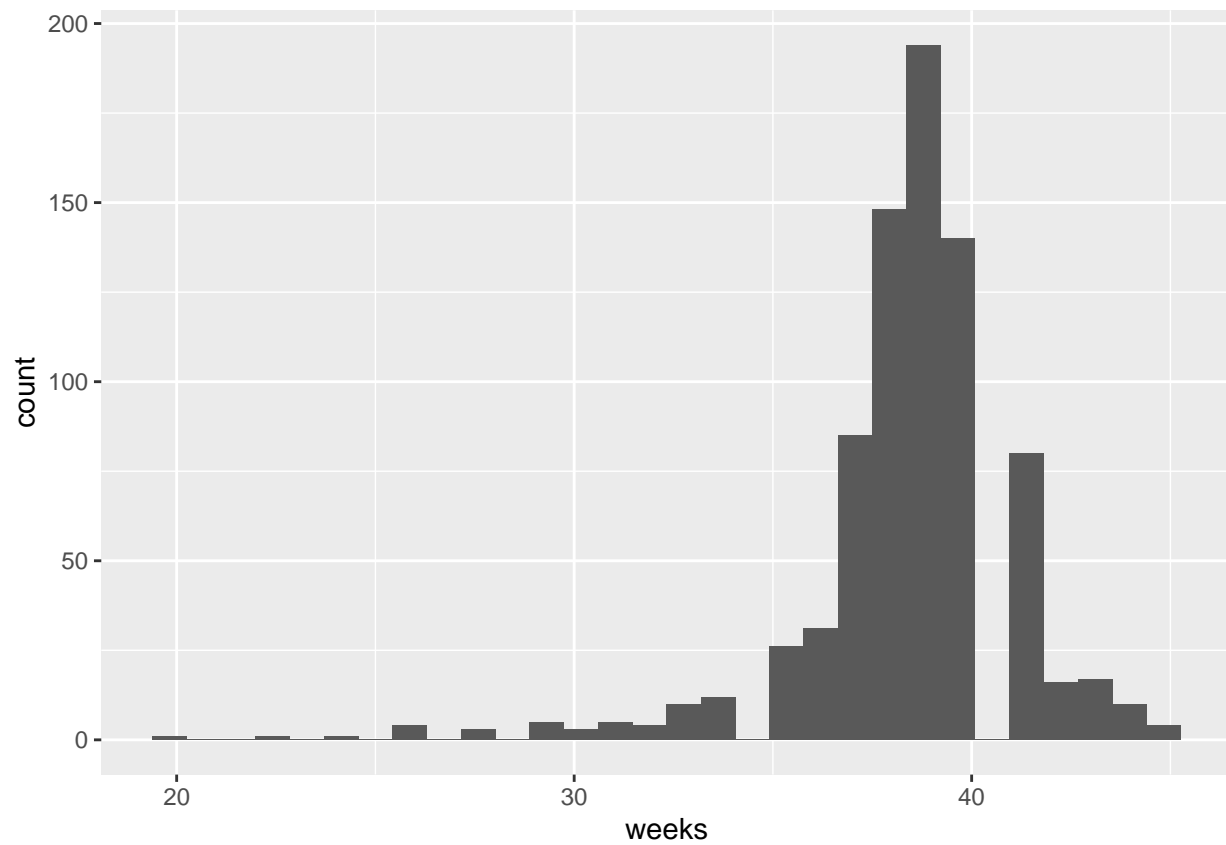
I can create graphical displays of data that highlight key features. I can combine multiple graphical displays or numerical summaries into an effective data product.

```

# histogram
ggplot(data = nc, aes(x = weeks))+
  geom_histogram()

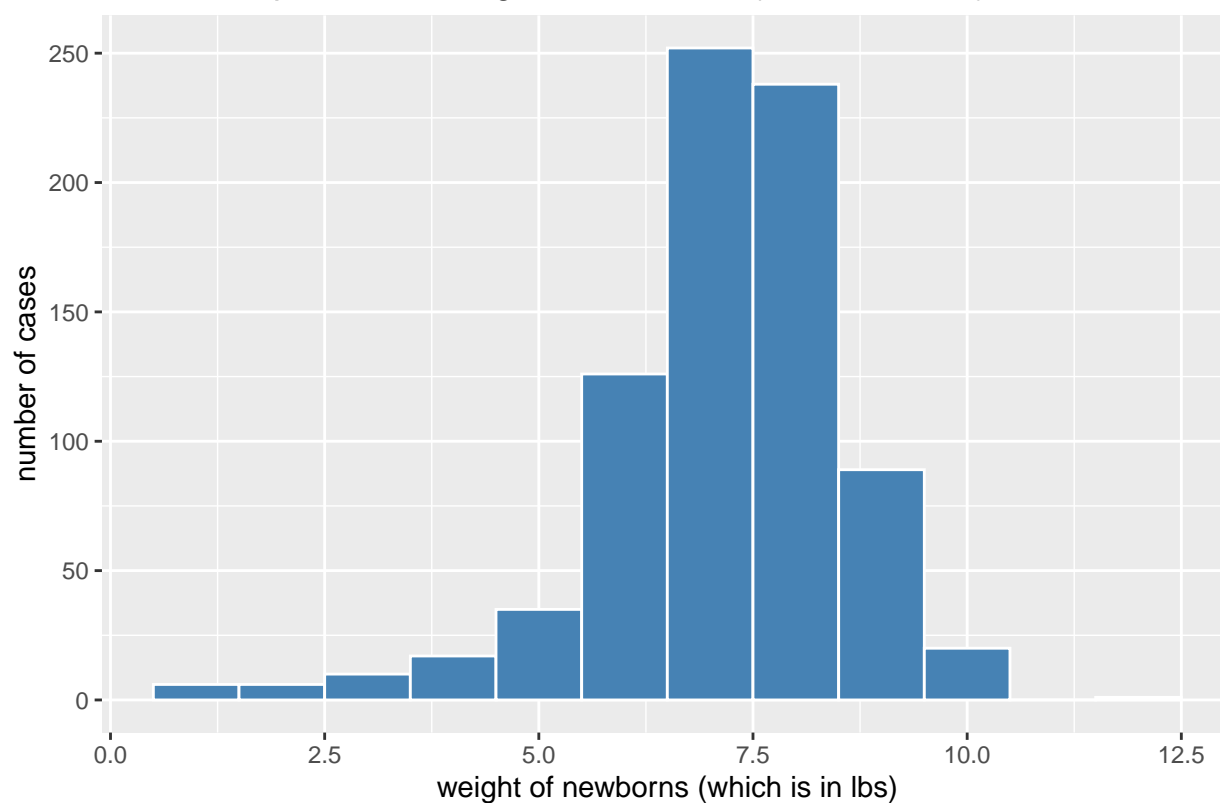
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

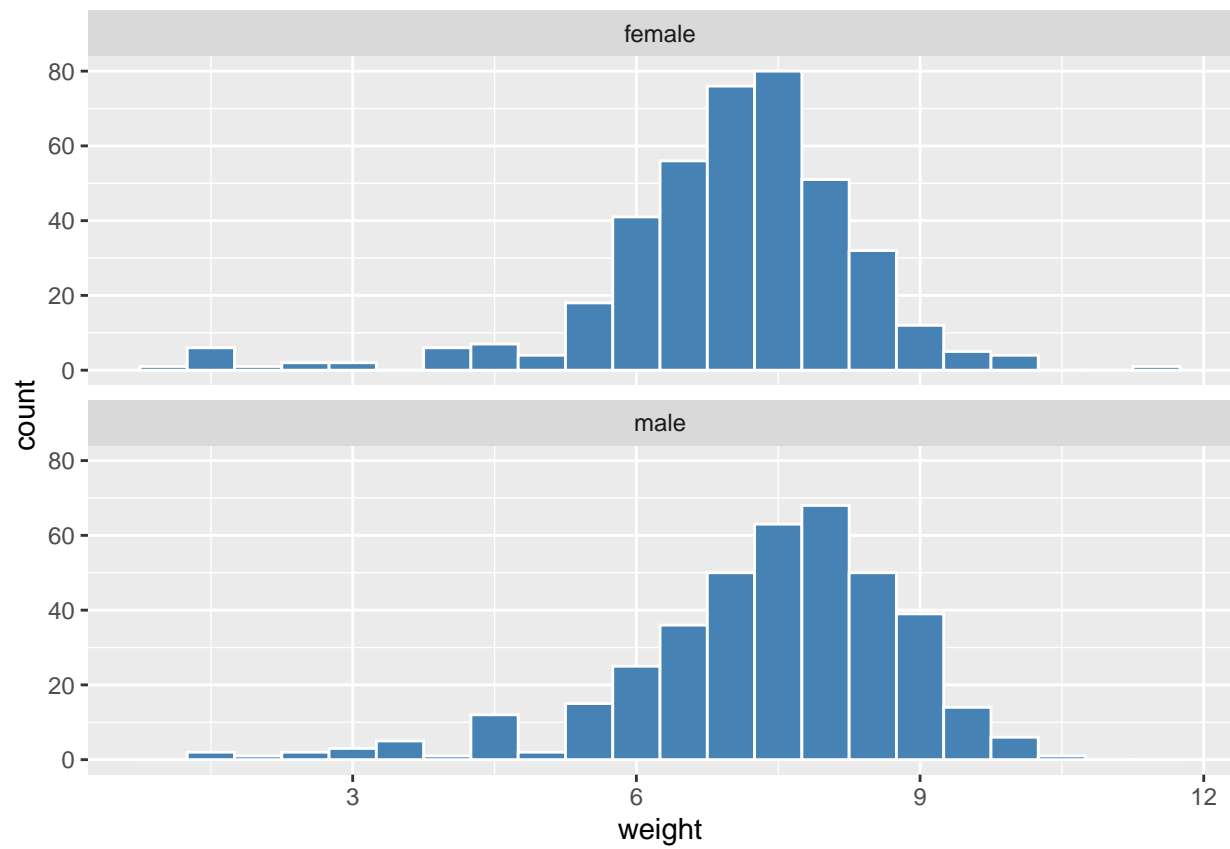


```
# collered and labeled histogram
ggplot(data = nc, aes(x = weight))+
  geom_histogram(binwidth = 1, color = "white", fill = "steelblue")+
  labs(x = "weight of newborns (which is in lbs)", y = "number of cases",
        title = "Relationship between weight of newborns (which is in lbs) and number of cases")
```

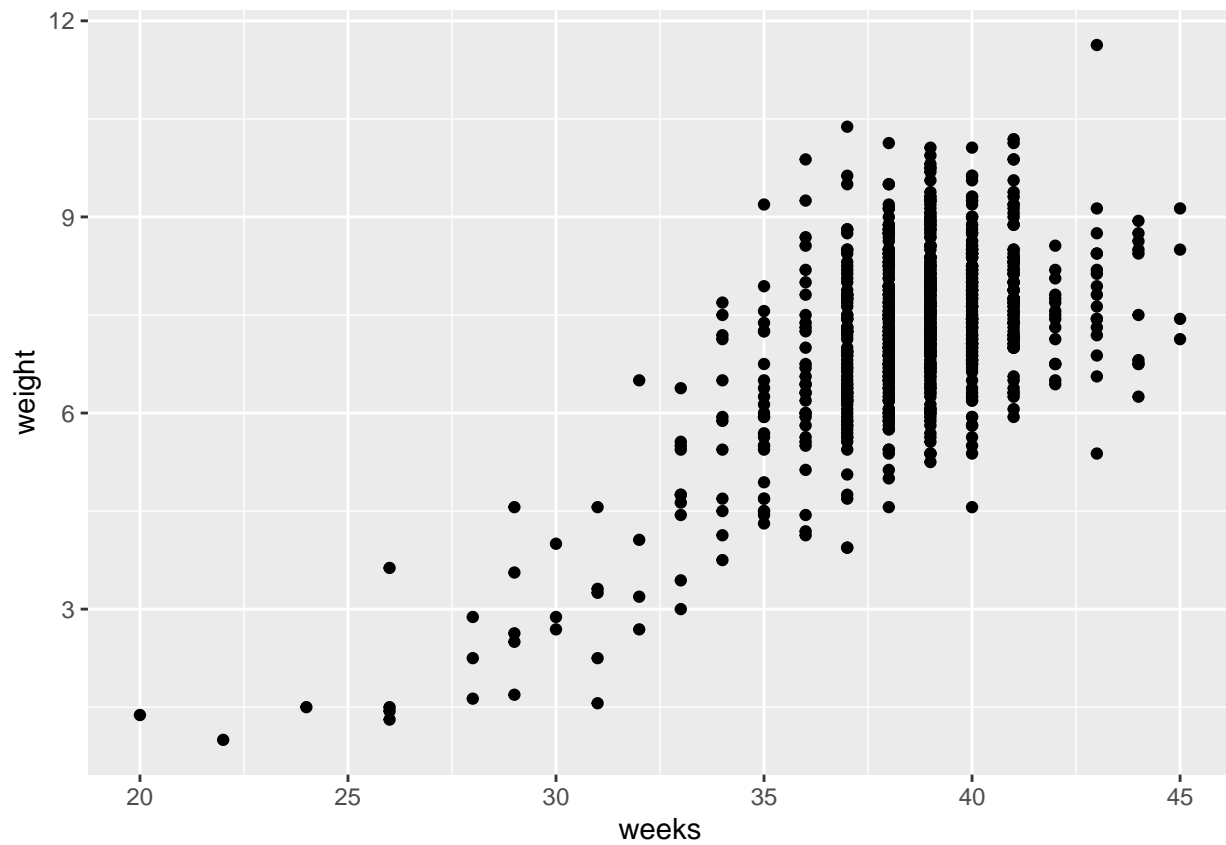
Relationship between weight of newborns (which is in lbs) and number of c



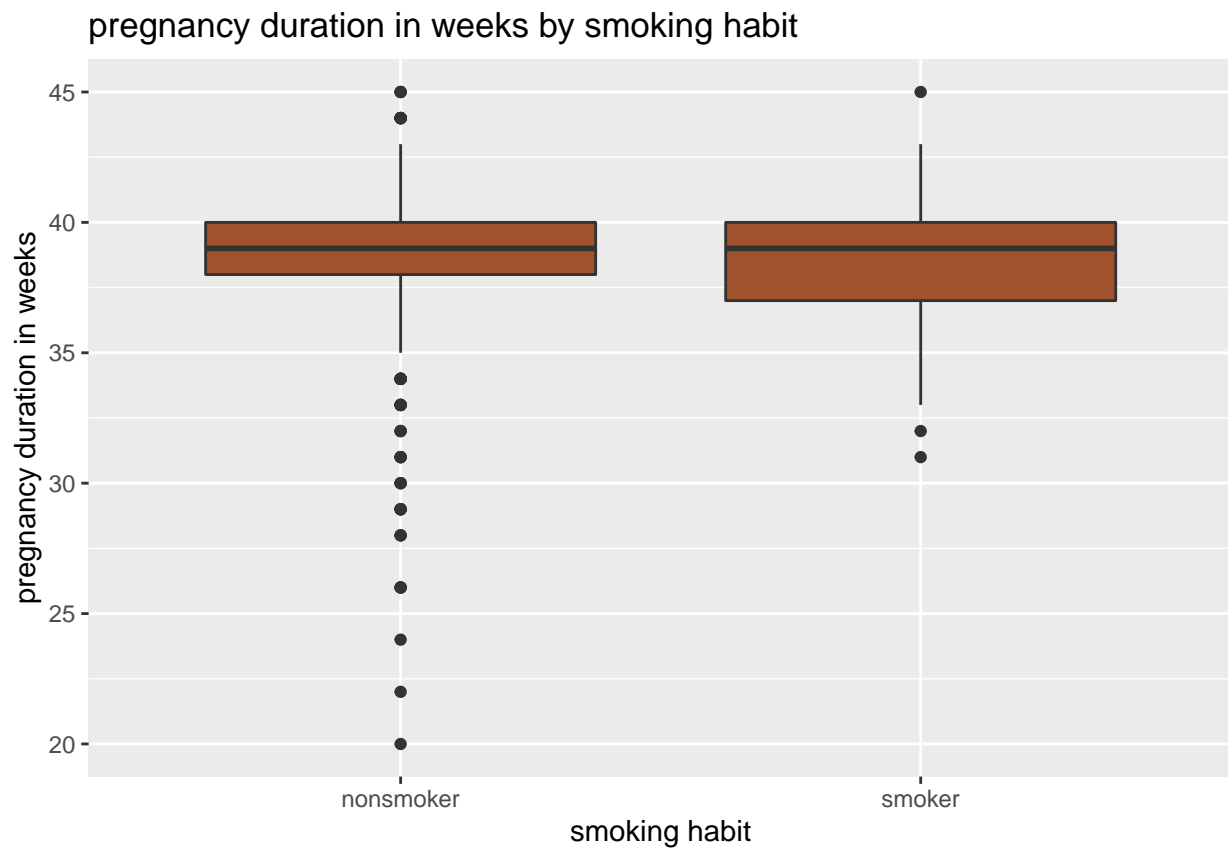
```
# two histograms in one plane  
ggplot(data = nc, aes(x = weight)) +  
  geom_histogram(binwidth = 0.5, color = "white", fill = "steelblue") +  
  facet_wrap(~ gender, ncol = 1)
```



```
# scatter plot
ggplot(data = nc, aes(x = weeks, y = weight)) +
  geom_point()
```

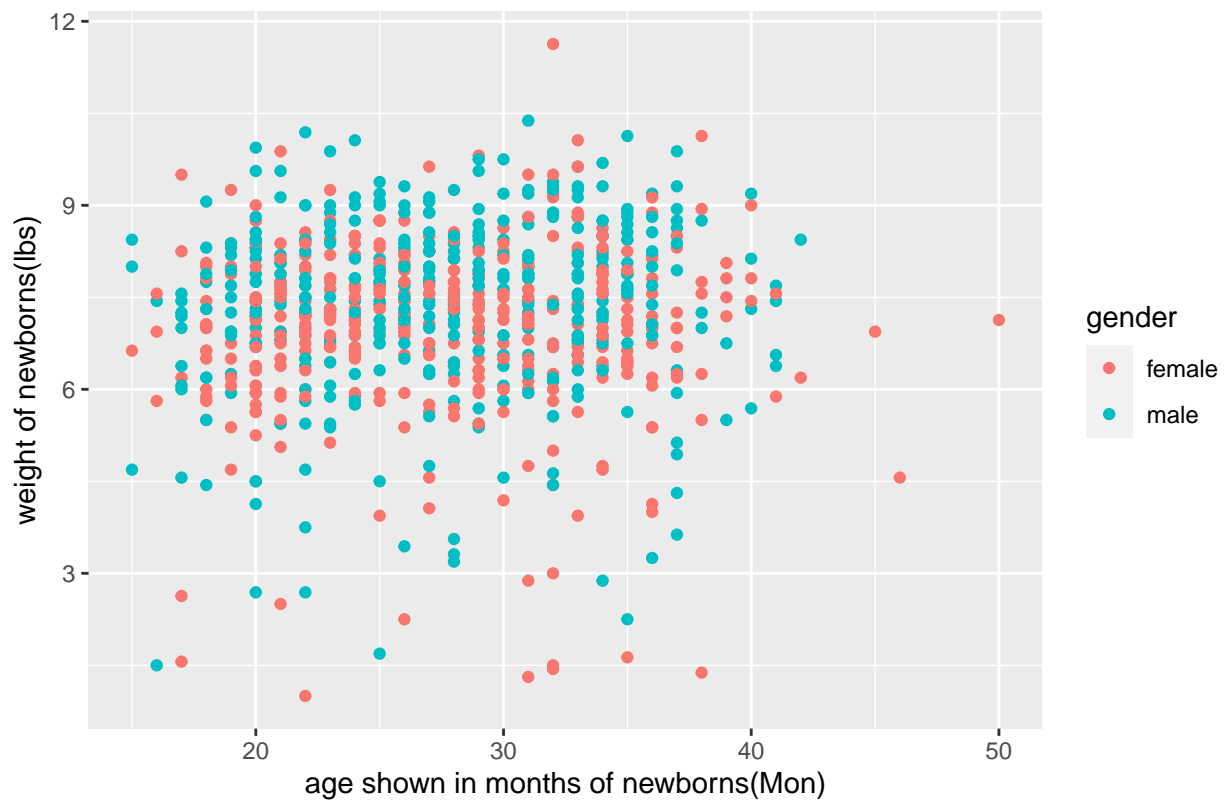


```
# box plot
ggplot(data = nc, aes(x = habit, y = weeks)) +
  geom_boxplot(fill = "sienna")+
  labs(x = "smoking habit", y = "pregnancy duration in weeks",
       title = "pregnancy duration in weeks by smoking habit")
```

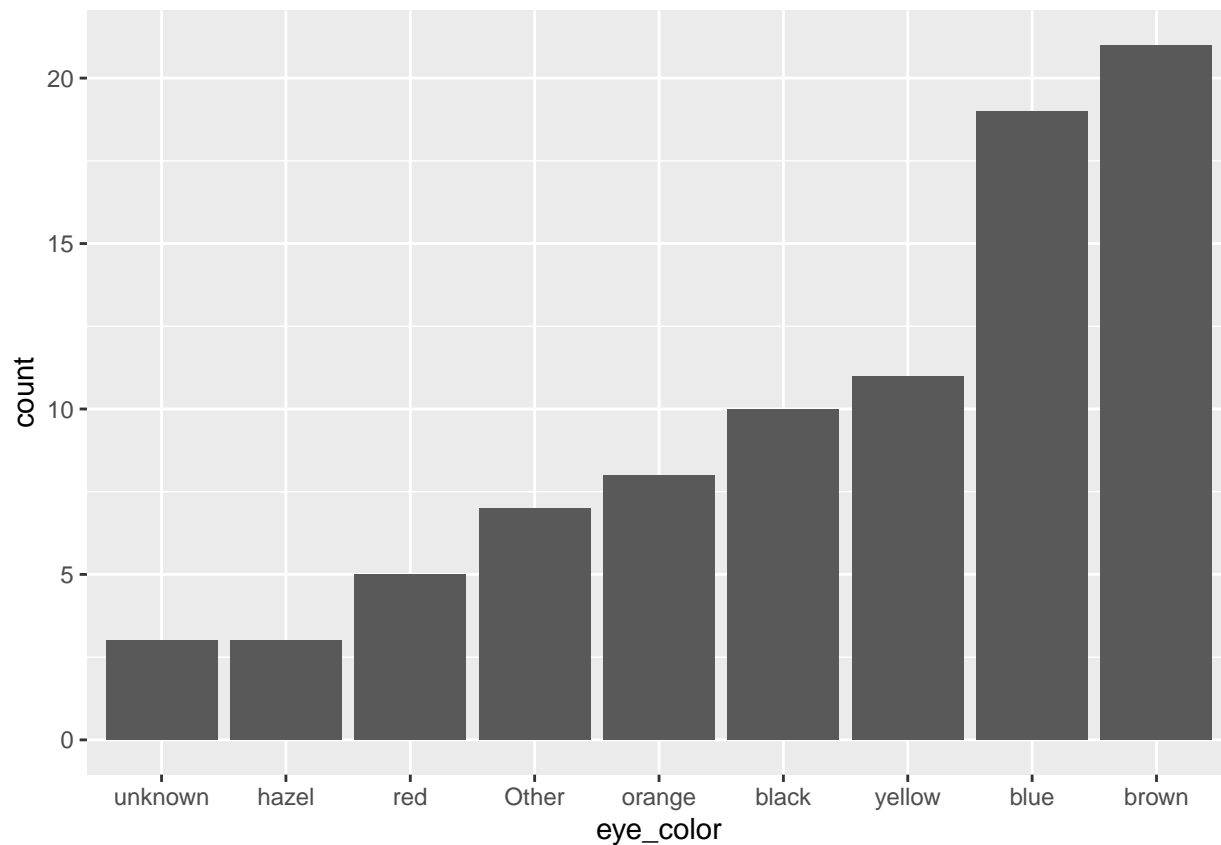


```
# adding color and labels to the plot
ggplot(data = nc, aes(x = mage, y = weight, color = gender))+
  geom_point() +
  labs(x = "age shown in months of newborns(Mon)", y = "weight of newborns(lbs)",
       title = "Relationship between age shown in months of newborns(Mon) and weight of newborns(lbs)")
```

Relationship between age shown in months of newborns(Mon) and weight o



```
# bar plot of the variable eye_color
people %>%
  mutate(eye_color = fct_lump(eye_color, prop = 0.03),
         eye_color = fct_infreq(eye_color),
         eye_color = fct_rev(eye_color)) %>%
  ggplot(mapping = aes(x = eye_color)) +
  geom_bar()
```

Creating my own functions

```
generate_n_samples <- function(n){
  tibble(
    normal_distribution = rnorm(n = n, mean = 50, sd = 10),
    exponential_distribution = rexp(n = n, rate = 0.2),
    binomial_distribution = rbinom(n = n, size = 100, prob = 0.25),
    uniform_distribution = runif(n = n, min = 10, max = 20)
  )
}
```

```
generate_n_samples(50)
```

```
## # A tibble: 50 x 4
##   normal_distributi~ exponential_distrib~ binomial_distribu~ uniform_distribut~
##           <dbl>           <dbl>           <int>           <dbl>
## 1           31.1           9.25             22           12.3
## 2           40.4           0.296            33           16.1
## 3           30.9           5.86             23           13.0
## 4           45.9           0.157            21           16.9
## 5           48.3           7.76             25           10.9
## 6           48.1           3.13             26           11.2
## 7           51.2           0.615            26           17.0
## 8           34.6           6.37             30           16.9
## 9           47.8          19.6             24           10.9
## 10          25.4           6.78             21           13.9
## # ... with 40 more rows
```

```
pow <- function(x, y) {
  # function to print x raised to the power y
  result <- x^y
  print(paste(x,"raised to the power", y, "is", result))
}
```

```
pow(2,10)
```

```
## [1] "2 raised to the power 10 is 1024"
```

```
pow(5,2)
```

```
## [1] "5 raised to the power 2 is 25"
```

```
# this function will return whether a given number is positive, negative or zero
check <- function(x) {
  if (x > 0) {
    result <- "Positive"
  }
  else if (x < 0) {
    result <- "Negative"
  }
  else {
    result <- "Zero"
  }
  return(result)
}
```

```
check(1)
```

```
## [1] "Positive"
```

```
check(-10)
```

```
## [1] "Negative"
```

```
check(0)
```

```
## [1] "Zero"
```

```
g <- factor(c("a","b","a","b","a","b","a","b","a","b","a","b"))
v <- c(1,4,1,4,1,4,2,8,2,8,2,8)
d <- data.frame(g,v)
d$cs <- ave(v, g, FUN=cumsum)
d
```

```
##      g v cs
## 1  a 1  1
## 2  b 4  4
## 3  a 1  2
## 4  b 4  8
## 5  a 1  3
## 6  b 4 12
## 7  a 2  5
## 8  b 8 20
## 9  a 2  7
## 10 b 8 28
## 11 a 2  9
```

```
## 12 b 8 36
```

Iterations

```
# using The repeat() Statement
Newton <- function(n, j=2, x=1) {
  # Use Newton's method to find the positive, real jth root of n,
  # where the default is to find the square root of n or j = 2.
  # x is a seed value to start the search.

  old.x <- x
  repeat {
    # Update x
    new.x <- old.x - ((old.x^j - n)) / (j * old.x^(j - 1))
    # Compute relative error as a 2-norm.
    conv <- sum((new.x - old.x)^2 / old.x^2)
    # Exit test with return() statement
    if(conv < 1e-10) return(old.x)
    # Save iteration result
    old.x <- new.x
  }
}
Newton(500, 2, 4)
```

```
## [1] 22.36068
```

```
# using The while() Statement
bit.string <- function(n) {
  tmp.string <- numeric(50)
  i <- 0
  while(n > 0) {
    tmp.string[50 - i] <- n %% 2 # modula
    n <- n %/% 2 # integer divide
    i <- i + 1
  }

  first.one <- match(1, tmp.string)
  return(tmp.string[first.one:50])
}

# Test the function
bit.string(1)
```

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

```
bit.string(2)
```

```
## [1] 1 0
```

```
bit.string(3)
```

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

```
bit.string(4)
```

```
## [1] 1 0 0
```

```
bit.string(5)
```

```
## [1] 1 0 1

# using The for() Statement
#creating a matrix using for()
x <- matrix(0, nrow = 3, ncol = 4)
for(i in 1:3) {
  for(j in 1:4) {
    x[i,j] <- i+j
  }
}
x

##      [,1] [,2] [,3] [,4]
## [1,]    2    3    4    5
## [2,]    3    4    5    6
## [3,]    4    5    6    7

# Use for() to create vector of cumulative sums
silly.csum <- function(N) {
  s <- vector("numeric", N)
  for (i in 1:N){
    if(i == 1) s[i] <- 1
    else s[i] <- s[i-1] + i;
  }
  return(s)
}

# Test the function
silly.csum(10)

## [1] 1 3 6 10 15 21 28 36 45 55

# Speed test1
system.time(silly.csum(1e7))

##      user  system elapsed
##  0.886   0.052   0.941

# speed test2
system.time(cumsum(as.numeric(1:1e7)))

##      user  system elapsed
##  0.064   0.085   0.149
```