

# tidying

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## packages

```
# install.packages ('gapminder')
library(gapminder)
library(tidyverse)

## -- Attaching packages ----- tidyverse 1.3.2 --
## v ggplot2 3.3.6      v purrr   0.3.4
## v tibble  3.1.8      v dplyr  1.0.10
## v tidyr   1.2.1      v stringr 1.4.1
## v readr   2.1.2      v forcats 0.5.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
```

## shortcuts

alt+- will add <-  
shift+ctrl+c to add # infront of a line  
'—' for a header, so it is easy to navigate through the script  
command +shift + m  
ctrl+alt+i for new code chunk # syntax Plain text  
End a line with two spaces to start a new paragraph.  
*italics* and *italics*  
**bold** and **bold**  
superscript<sup>2</sup>  
~strikethrough  
link to rstudio

## logical operations

```
1==1 # equality
1!=3 #unequal
13<14 #13 smaller than 14
14>13 #14 bigger than 13
12>=0 #12 greater or equal to zero
12<=3 #12 smaller or equal to zero
```

## creating data.frame

family

```
name <- c('saneesh', 'sanusha', 'appu', 'kishan')
weight <- c(63, 48, 20, NA)
height <- c(164, 150, NA, 75)
family <- data.frame(name, weight, height)
family %>% as_tibble()
```

```
## # A tibble: 4 x 3
##   name    weight height
##   <chr>    <dbl> <dbl>
## 1 saneesh     63    164
## 2 sanusha     48    150
## 3 appu        20     NA
## 4 kishan      NA     75
```

```
library(tidyverse)
data <- data.frame(sex=c(rep('female', 10), rep('male', 8)),
                  score=c(rnorm(n= 10, mean = 7.56, sd = 1.978), rnorm(n= 8, mean=7.75, sd= 1.631)))
data
```

data frame with unequal values 10 and 8

```
##      sex      score
## 1 female 6.9540600
## 2 female 9.4259315
## 3 female 9.1106210
## 4 female 7.2967252
## 5 female 0.5668015
## 6 female 7.5257822
## 7 female 6.2697928
## 8 female 6.9339602
## 9 female 6.1535163
## 10 female 3.4910928
## 11 male 8.8765558
## 12 male 4.7396814
## 13 male 7.0524847
## 14 male 8.5663857
## 15 male 8.7699144
## 16 male 7.1089717
## 17 male 6.4231238
## 18 male 8.3191404
```

```
data %>% group_by(sex) %>%
  summarise(score= n()) %>%
  mutate(freq=score/sum(score)*100)
```

```
## # A tibble: 2 x 3
##   sex    score freq
##   <chr> <int> <dbl>
## 1 female    10  55.6
## 2 male      8  44.4
```

## is.na

```
# identify location of NAs in vector
which(is.na(family))
```

```
## [1] 8 11
```

```
colSums(is.na(family))
```

```
##   name weight height
##     0      1      1
```

## replace na

```
mat <- matrix(sample(c(NA, 1:5), 50, replace = TRUE), 5)
df <- as.data.frame(mat)
df %>% replace(is.na(.), 0)%>% view()
```

## clean names

```
# install.packages('janitor')
library(janitor)
```

```
##
## Attaching package: 'janitor'
```

```
## The following objects are masked from 'package:stats':
##
##   chisq.test, fisher.test
```

```
id <- (c(1,1,2,2,3,3))
Country <- c('Angola', 'Angola', 'Botswana', 'Botswana', 'Zimbabwe', 'Zimbabwe')
year <- c('2006', '2007', '2008', '2009', '2010', '2006')
bank.ratio <- c(24,25,38,34,42,49)
Reserve.ratio <- c(77,59,64,65,57,86)
broad.money <- c(163,188,317,361,150,288)
```

```
bank <- data.frame(id, Country, year, bank.ratio, Reserve.ratio,broad.money)

bank %>% view()
as_tibble()
```

## Warning: The `x` argument of `as\_tibble()` can't be missing as of tibble 3.0.0.

## # A tibble: 0 x 0

```
bank <- bank %>% clean_names() # replaced . with _

glimpse(bank)
```

```
## Rows: 6
## Columns: 6
## $ id          <dbl> 1, 1, 2, 2, 3, 3
## $ country     <chr> "Angola", "Angola", "Botswana", "Botswana", "Zimbabwe", ~
## $ year        <chr> "2006", "2007", "2008", "2009", "2010", "2006"
## $ bank_ratio  <dbl> 24, 25, 38, 34, 42, 49
## $ reserve_ratio <dbl> 77, 59, 64, 65, 57, 86
## $ broad_money <dbl> 163, 188, 317, 361, 150, 288
```

```
bank <- bank %>% clean_names() # replaced . with _
```

**filter** bank data frame below such that it retains a country if a given id is satisfied e.g. filtering a data frame that has countries with id 1 and 2 only

```
bank %>%
  filter(id%in% c(1,2)) %>%
  as_tibble()
```

```
## # A tibble: 4 x 6
##       id country  year bank_ratio reserve_ratio broad_money
##   <dbl> <chr>   <chr>      <dbl>         <dbl>         <dbl>
## 1     1  Angola  2006         24           77           163
## 2     1  Angola  2007         25           59           188
## 3     2 Botswana 2008         38           64           317
## 4     2 Botswana 2009         34           65           361
```

**summarise** fund available with each countries

```
bank %>%
  group_by(country) %>%
  summarise(fund=sum(broad_money)) %>%
  as_tibble()
```

```
## # A tibble: 3 x 2
##   country  fund
##   <chr>   <dbl>
## 1 Angola    351
## 2 Botswana  678
## 3 Zimbabwe  438
```

## count/ frequency

```
mtcars %>%  
  count(am) %>%  
  as_tibble()
```

```
## # A tibble: 2 x 2  
##       am       n  
##   <dbl> <int>  
## 1     0     19  
## 2     1     13
```

```
mtcars %>%  
  count(gear)
```

```
##   gear  n  
## 1    3 15  
## 2    4 12  
## 3    5  5
```

## rename column

column: new name= old name

```
iris %>%  
  rename(sep.len=Sepal.Length)
```

```
##       sep.len Sepal.Width Petal.Length Petal.Width Species  
## 1      5.1      3.5      1.4      0.2    setosa  
## 2      4.9      3.0      1.4      0.2    setosa  
## 3      4.7      3.2      1.3      0.2    setosa  
## 4      4.6      3.1      1.5      0.2    setosa  
## 5      5.0      3.6      1.4      0.2    setosa  
## 6      5.4      3.9      1.7      0.4    setosa  
## 7      4.6      3.4      1.4      0.3    setosa  
## 8      5.0      3.4      1.5      0.2    setosa  
## 9      4.4      2.9      1.4      0.2    setosa  
## 10     4.9      3.1      1.5      0.1    setosa  
## 11     5.4      3.7      1.5      0.2    setosa  
## 12     4.8      3.4      1.6      0.2    setosa  
## 13     4.8      3.0      1.4      0.1    setosa  
## 14     4.3      3.0      1.1      0.1    setosa  
## 15     5.8      4.0      1.2      0.2    setosa  
## 16     5.7      4.4      1.5      0.4    setosa  
## 17     5.4      3.9      1.3      0.4    setosa  
## 18     5.1      3.5      1.4      0.3    setosa  
## 19     5.7      3.8      1.7      0.3    setosa  
## 20     5.1      3.8      1.5      0.3    setosa  
## 21     5.4      3.4      1.7      0.2    setosa
```

## 22	5.1	3.7	1.5	0.4	setosa
## 23	4.6	3.6	1.0	0.2	setosa
## 24	5.1	3.3	1.7	0.5	setosa
## 25	4.8	3.4	1.9	0.2	setosa
## 26	5.0	3.0	1.6	0.2	setosa
## 27	5.0	3.4	1.6	0.4	setosa
## 28	5.2	3.5	1.5	0.2	setosa
## 29	5.2	3.4	1.4	0.2	setosa
## 30	4.7	3.2	1.6	0.2	setosa
## 31	4.8	3.1	1.6	0.2	setosa
## 32	5.4	3.4	1.5	0.4	setosa
## 33	5.2	4.1	1.5	0.1	setosa
## 34	5.5	4.2	1.4	0.2	setosa
## 35	4.9	3.1	1.5	0.2	setosa
## 36	5.0	3.2	1.2	0.2	setosa
## 37	5.5	3.5	1.3	0.2	setosa
## 38	4.9	3.6	1.4	0.1	setosa
## 39	4.4	3.0	1.3	0.2	setosa
## 40	5.1	3.4	1.5	0.2	setosa
## 41	5.0	3.5	1.3	0.3	setosa
## 42	4.5	2.3	1.3	0.3	setosa
## 43	4.4	3.2	1.3	0.2	setosa
## 44	5.0	3.5	1.6	0.6	setosa
## 45	5.1	3.8	1.9	0.4	setosa
## 46	4.8	3.0	1.4	0.3	setosa
## 47	5.1	3.8	1.6	0.2	setosa
## 48	4.6	3.2	1.4	0.2	setosa
## 49	5.3	3.7	1.5	0.2	setosa
## 50	5.0	3.3	1.4	0.2	setosa
## 51	7.0	3.2	4.7	1.4	versicolor
## 52	6.4	3.2	4.5	1.5	versicolor
## 53	6.9	3.1	4.9	1.5	versicolor
## 54	5.5	2.3	4.0	1.3	versicolor
## 55	6.5	2.8	4.6	1.5	versicolor
## 56	5.7	2.8	4.5	1.3	versicolor
## 57	6.3	3.3	4.7	1.6	versicolor
## 58	4.9	2.4	3.3	1.0	versicolor
## 59	6.6	2.9	4.6	1.3	versicolor
## 60	5.2	2.7	3.9	1.4	versicolor
## 61	5.0	2.0	3.5	1.0	versicolor
## 62	5.9	3.0	4.2	1.5	versicolor
## 63	6.0	2.2	4.0	1.0	versicolor
## 64	6.1	2.9	4.7	1.4	versicolor
## 65	5.6	2.9	3.6	1.3	versicolor
## 66	6.7	3.1	4.4	1.4	versicolor
## 67	5.6	3.0	4.5	1.5	versicolor
## 68	5.8	2.7	4.1	1.0	versicolor
## 69	6.2	2.2	4.5	1.5	versicolor
## 70	5.6	2.5	3.9	1.1	versicolor
## 71	5.9	3.2	4.8	1.8	versicolor
## 72	6.1	2.8	4.0	1.3	versicolor
## 73	6.3	2.5	4.9	1.5	versicolor
## 74	6.1	2.8	4.7	1.2	versicolor
## 75	6.4	2.9	4.3	1.3	versicolor

## 76	6.6	3.0	4.4	1.4 versicolor
## 77	6.8	2.8	4.8	1.4 versicolor
## 78	6.7	3.0	5.0	1.7 versicolor
## 79	6.0	2.9	4.5	1.5 versicolor
## 80	5.7	2.6	3.5	1.0 versicolor
## 81	5.5	2.4	3.8	1.1 versicolor
## 82	5.5	2.4	3.7	1.0 versicolor
## 83	5.8	2.7	3.9	1.2 versicolor
## 84	6.0	2.7	5.1	1.6 versicolor
## 85	5.4	3.0	4.5	1.5 versicolor
## 86	6.0	3.4	4.5	1.6 versicolor
## 87	6.7	3.1	4.7	1.5 versicolor
## 88	6.3	2.3	4.4	1.3 versicolor
## 89	5.6	3.0	4.1	1.3 versicolor
## 90	5.5	2.5	4.0	1.3 versicolor
## 91	5.5	2.6	4.4	1.2 versicolor
## 92	6.1	3.0	4.6	1.4 versicolor
## 93	5.8	2.6	4.0	1.2 versicolor
## 94	5.0	2.3	3.3	1.0 versicolor
## 95	5.6	2.7	4.2	1.3 versicolor
## 96	5.7	3.0	4.2	1.2 versicolor
## 97	5.7	2.9	4.2	1.3 versicolor
## 98	6.2	2.9	4.3	1.3 versicolor
## 99	5.1	2.5	3.0	1.1 versicolor
## 100	5.7	2.8	4.1	1.3 versicolor
## 101	6.3	3.3	6.0	2.5 virginica
## 102	5.8	2.7	5.1	1.9 virginica
## 103	7.1	3.0	5.9	2.1 virginica
## 104	6.3	2.9	5.6	1.8 virginica
## 105	6.5	3.0	5.8	2.2 virginica
## 106	7.6	3.0	6.6	2.1 virginica
## 107	4.9	2.5	4.5	1.7 virginica
## 108	7.3	2.9	6.3	1.8 virginica
## 109	6.7	2.5	5.8	1.8 virginica
## 110	7.2	3.6	6.1	2.5 virginica
## 111	6.5	3.2	5.1	2.0 virginica
## 112	6.4	2.7	5.3	1.9 virginica
## 113	6.8	3.0	5.5	2.1 virginica
## 114	5.7	2.5	5.0	2.0 virginica
## 115	5.8	2.8	5.1	2.4 virginica
## 116	6.4	3.2	5.3	2.3 virginica
## 117	6.5	3.0	5.5	1.8 virginica
## 118	7.7	3.8	6.7	2.2 virginica
## 119	7.7	2.6	6.9	2.3 virginica
## 120	6.0	2.2	5.0	1.5 virginica
## 121	6.9	3.2	5.7	2.3 virginica
## 122	5.6	2.8	4.9	2.0 virginica
## 123	7.7	2.8	6.7	2.0 virginica
## 124	6.3	2.7	4.9	1.8 virginica
## 125	6.7	3.3	5.7	2.1 virginica
## 126	7.2	3.2	6.0	1.8 virginica
## 127	6.2	2.8	4.8	1.8 virginica
## 128	6.1	3.0	4.9	1.8 virginica
## 129	6.4	2.8	5.6	2.1 virginica

```
## 130      7.2      3.0      5.8      1.6 virginica
## 131      7.4      2.8      6.1      1.9 virginica
## 132      7.9      3.8      6.4      2.0 virginica
## 133      6.4      2.8      5.6      2.2 virginica
## 134      6.3      2.8      5.1      1.5 virginica
## 135      6.1      2.6      5.6      1.4 virginica
## 136      7.7      3.0      6.1      2.3 virginica
## 137      6.3      3.4      5.6      2.4 virginica
## 138      6.4      3.1      5.5      1.8 virginica
## 139      6.0      3.0      4.8      1.8 virginica
## 140      6.9      3.1      5.4      2.1 virginica
## 141      6.7      3.1      5.6      2.4 virginica
## 142      6.9      3.1      5.1      2.3 virginica
## 143      5.8      2.7      5.1      1.9 virginica
## 144      6.8      3.2      5.9      2.3 virginica
## 145      6.7      3.3      5.7      2.5 virginica
## 146      6.7      3.0      5.2      2.3 virginica
## 147      6.3      2.5      5.0      1.9 virginica
## 148      6.5      3.0      5.2      2.0 virginica
## 149      6.2      3.4      5.4      2.3 virginica
## 150      5.9      3.0      5.1      1.8 virginica
```

## gapminder

```
iris %>% as_tibble()
```

```
## # A tibble: 150 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##   <dbl>         <dbl>         <dbl>         <dbl> <fct>
## 1           5.1           3.5           1.4           0.2 setosa
## 2           4.9           3           1.4           0.2 setosa
## 3           4.7           3.2           1.3           0.2 setosa
## 4           4.6           3.1           1.5           0.2 setosa
## 5           5           3.6           1.4           0.2 setosa
## 6           5.4           3.9           1.7           0.4 setosa
## 7           4.6           3.4           1.4           0.3 setosa
## 8           5           3.4           1.5           0.2 setosa
## 9           4.4           2.9           1.4           0.2 setosa
## 10          4.9           3.1           1.5           0.1 setosa
## # ... with 140 more rows
```

```
summary(gapminder)
```

```
##           country           continent           year           lifeExp
## Afghanistan: 12 Africa :624 Min. :1952 Min. :23.60
## Albania : 12 Americas:300 1st Qu.:1966 1st Qu.:48.20
## Algeria : 12 Asia :396 Median :1980 Median :60.71
## Angola : 12 Europe :360 Mean :1980 Mean :59.47
## Argentina : 12 Oceania : 24 3rd Qu.:1993 3rd Qu.:70.85
## Australia : 12 Max. :2007 Max. :82.60
```



```
## (Other)      :1632
##      pop      gdpPercap
## Min.      :6.001e+04   Min.      : 241.2
## 1st Qu.:2.794e+06   1st Qu.: 1202.1
## Median :7.024e+06   Median : 3531.8
## Mean      :2.960e+07   Mean      : 7215.3
## 3rd Qu.:1.959e+07   3rd Qu.: 9325.5
## Max.      :1.319e+09   Max.      :113523.1
##
```

```
str(gapminder)
```

```
## tibble [1,704 x 6] (S3: tbl_df/tbl/data.frame)
## $ country : Factor w/ 142 levels "Afghanistan",...: 1 1 1 1 1 1 1 1 1 ...
## $ continent: Factor w/ 5 levels "Africa","Americas",...: 3 3 3 3 3 3 3 3 3 ...
## $ year      : int [1:1704] 1952 1957 1962 1967 1972 1977 1982 1987 1992 1997 ...
## $ lifeExp   : num [1:1704] 28.8 30.3 32 34 36.1 ...
## $ pop       : int [1:1704] 8425333 9240934 10267083 11537966 13079460 14880372 12881816 13867957 163...
## $ gdpPercap: num [1:1704] 779 821 853 836 740 ...
```

## recode observation

change name of observation— mutate (variable=recode (variable, ‘old name’=‘new name’)))

```
gapminder %>%
  mutate(country=recode(country, 'India'='IND' )) %>%
  filter(country=='IND')
```

```
## # A tibble: 12 x 6
##   country continent  year lifeExp      pop gdpPercap
##   <fct>    <fct>    <int>  <dbl>    <int>  <dbl>
## 1 IND      Asia      1952   37.4  372000000    547.
## 2 IND      Asia      1957   40.2  409000000    590.
## 3 IND      Asia      1962   43.6  454000000    658.
## 4 IND      Asia      1967   47.2  506000000    701.
## 5 IND      Asia      1972   50.7  567000000    724.
## 6 IND      Asia      1977   54.2  634000000    813.
## 7 IND      Asia      1982   56.6  708000000    856.
## 8 IND      Asia      1987   58.6  788000000    977.
## 9 IND      Asia      1992   60.2  872000000   1164.
## 10 IND     Asia      1997   61.8  959000000   1459.
## 11 IND     Asia      2002   62.9 1034172547   1747.
## 12 IND     Asia      2007   64.7 1110396331   2452.
```

## select

```
gapminder %>%
  select(year, country, gdpPercap)
```

```
## # A tibble: 1,704 x 3
##   year country      gdpPercap
##   <int> <fct>      <dbl>
## 1  1952 Afghanistan    779.
## 2  1957 Afghanistan    821.
## 3  1962 Afghanistan    853.
## 4  1967 Afghanistan    836.
## 5  1972 Afghanistan    740.
## 6  1977 Afghanistan    786.
## 7  1982 Afghanistan    978.
## 8  1987 Afghanistan    852.
## 9  1992 Afghanistan    649.
## 10 1997 Afghanistan    635.
## # ... with 1,694 more rows
```

```
msleep %>% select(starts_with("sleep"))
```

```
## # A tibble: 83 x 3
##   sleep_total sleep_rem sleep_cycle
##   <dbl>      <dbl>      <dbl>
## 1      12.1      NA        NA
## 2       17       1.8        NA
## 3      14.4       2.4        NA
## 4      14.9       2.3      0.133
## 5        4       0.7      0.667
## 6      14.4       2.2      0.767
## 7       8.7       1.4      0.383
## 8        7       NA        NA
## 9      10.1       2.9      0.333
## 10       3       NA        NA
## # ... with 73 more rows
```

## filter

```
gapminder %>%
  select(year, country, lifeExp) %>%
  filter(country=="Eritrea", year>1950)
```

```
## # A tibble: 12 x 3
##   year country lifeExp
##   <int> <fct>      <dbl>
## 1  1952 Eritrea    35.9
## 2  1957 Eritrea    38.0
## 3  1962 Eritrea    40.2
## 4  1967 Eritrea    42.2
## 5  1972 Eritrea    44.1
## 6  1977 Eritrea    44.5
## 7  1982 Eritrea    43.9
## 8  1987 Eritrea    46.5
## 9  1992 Eritrea    50.0
```

```
## 10 1997 Eritrea 53.4
## 11 2002 Eritrea 55.2
## 12 2007 Eritrea 58.0
```

```
gapminder %>% filter(country=="Canada") %>% head(3) # from gapminder data filter country Canada and show
```

```
## # A tibble: 3 x 6
##   country continent year lifeExp      pop gdpPercap
##   <fct>    <fct>    <int>   <dbl>    <int>    <dbl>
## 1 Canada  Americas  1952   68.8 14785584  11367.
## 2 Canada  Americas  1957   70.0 17010154  12490.
## 3 Canada  Americas  1962   71.3 18985849  13462.
```

```
gapminder %>% filter(country!="Oman") %>% head(3) # from gapminder data filter all the other countries
```

```
## # A tibble: 3 x 6
##   country      continent year lifeExp      pop gdpPercap
##   <fct>        <fct>    <int>   <dbl>    <int>    <dbl>
## 1 Afghanistan Asia      1952   28.8  8425333    779.
## 2 Afghanistan Asia      1957   30.3  9240934    821.
## 3 Afghanistan Asia      1962   32.0 10267083    853.
```

multiple conditions

```
gapminder %>%
  filter(country=="Oman" &
         year>1980 &
         year<=2000) %>% head(4)
```

```
## # A tibble: 4 x 6
##   country continent year lifeExp      pop gdpPercap
##   <fct>    <fct>    <int>   <dbl>    <int>    <dbl>
## 1 Oman     Asia      1982   62.7 1301048  12955.
## 2 Oman     Asia      1987   67.7 1593882  18115.
## 3 Oman     Asia      1992   71.2 1915208  18617.
## 4 Oman     Asia      1997   72.5 2283635  19702.
```

```
gapminder %>%
  select(country, year) %>%
  filter(year>=1980, country=="India" |
         country=="Oman" |
         country=="Canada")
```

```
## # A tibble: 18 x 2
##   country year
##   <fct>   <int>
## 1 Canada  1982
## 2 Canada  1987
## 3 Canada  1992
## 4 Canada  1997
## 5 Canada  2002
```

```
## 6 Canada 2007
## 7 India 1982
## 8 India 1987
## 9 India 1992
## 10 India 1997
## 11 India 2002
## 12 India 2007
## 13 Oman 1982
## 14 Oman 1987
## 15 Oman 1992
## 16 Oman 1997
## 17 Oman 2002
## 18 Oman 2007
```

```
gapminder %>% filter(country!="Oman") %>% head(3) # from gapminder data filter all the other countries
```

```
## # A tibble: 3 x 6
##   country    continent year lifeExp    pop gdpPercap
##   <fct>      <fct>    <int>   <dbl>   <int>   <dbl>
## 1 Afghanistan Asia      1952    28.8  8425333    779.
## 2 Afghanistan Asia      1957    30.3  9240934    821.
## 3 Afghanistan Asia      1962    32.0 10267083    853.
```

## drop

```
gapminder %>%
  select(-year,-pop) %>%
  head(5)
```

```
## # A tibble: 5 x 4
##   country    continent lifeExp gdpPercap
##   <fct>      <fct>      <dbl>   <dbl>
## 1 Afghanistan Asia      28.8    779.
## 2 Afghanistan Asia      30.3    821.
## 3 Afghanistan Asia      32.0    853.
## 4 Afghanistan Asia      34.0    836.
## 5 Afghanistan Asia      36.1    740.
```

## group by & summarise

```
gapminder %>%
  filter(year==2007) %>%
  group_by(country) %>%
  summarise(meanLE=mean(lifeExp)) %>%
  arrange(meanLE,decreasing = TRUE)
```

```
## # A tibble: 142 x 2
```

```
##   country      meanLE
##   <fct>        <dbl>
## 1 Swaziland    39.6
## 2 Mozambique    42.1
## 3 Zambia        42.4
## 4 Sierra Leone 42.6
## 5 Lesotho       42.6
## 6 Angola        42.7
## 7 Zimbabwe      43.5
## 8 Afghanistan   43.8
## 9 Central African Republic 44.7
## 10 Liberia      45.7
## # ... with 132 more rows
```

```
gapminder %>%
  group_by(country) %>%
  summarise(minLE=min(lifeExp)) %>%
  arrange(minLE,decreasing=FALSE)
```

```
## # A tibble: 142 x 2
##   country      minLE
##   <fct>        <dbl>
## 1 Rwanda      23.6
## 2 Afghanistan 28.8
## 3 Gambia       30
## 4 Angola       30.0
## 5 Sierra Leone 30.3
## 6 Cambodia     31.2
## 7 Mozambique   31.3
## 8 Burkina Faso 32.0
## 9 Guinea-Bissau 32.5
## 10 Yemen, Rep. 32.5
## # ... with 132 more rows
```

grouped by continent, then summarise two things, first `n=n()` number of rows in which each continent are or the size of each group, then the mean of the mean of the `lifeExp` variable.

```
gapminder %>%
  group_by(continent) %>%
  summarise(n=n(),
            meanLife=mean(lifeExp))
```

```
## # A tibble: 5 x 3
##   continent      n meanLife
##   <fct>    <int>    <dbl>
## 1 Africa    624    48.9
## 2 Americas  300    64.7
## 3 Asia     396    60.1
## 4 Europe   360    71.9
## 5 Oceania   24     74.3
```

```
gapminder %>%
  group_by(continent) %>%
  summarise(PopConti=sum(pop))
```

```
## # A tibble: 5 x 2
##   continent    PopConti
##   <fct>         <dbl>
## 1 Africa      6187585961
## 2 Americas    7351438499
## 3 Asia        30507333901
## 4 Europe      6181115304
## 5 Oceania     212992136
```

```
pets <- data.frame(names=c(rep('saneesh', 3), rep('appu', 2), 'sanusha'),
  pet=c(rep('dog', 3), rep('cat', 2), 'tiger'), number=c(2,2,5,7,8,1),
  size=c(rep('medium', 2), rep('small', 3), 'big'))
```

```
pets
```

```
##   names    pet number  size
## 1 saneesh  dog      2 medium
## 2 saneesh  dog      2 medium
## 3 saneesh  dog      5 small
## 4 appu     cat      7 small
## 5 appu     cat      8 small
## 6 sanusha  tiger    1 big
```

```
library(tidyverse)
```

```
pets %>% group_by(pet, size) %>%
  summarise(totalpet= sum(number))
```

```
## `summarise()` has grouped output by 'pet'. You can override using the `.groups`
## argument.
```

```
## # A tibble: 4 x 3
## # Groups:   pet [3]
##   pet    size  totalpet
##   <chr> <chr>      <dbl>
## 1 cat   small        15
## 2 dog   medium         4
## 3 dog   small         5
## 4 tiger big          1
```

## summarise

```
library(tidyverse)
plot <- c(rep(1,2), rep(2,4), rep(3,3))
```

```
bird <- c('a','b', 'a','b', 'c', 'd', 'a', 'b', 'c')
area <- c(rep(10,2), rep(5,4), rep(15,3))

birdlist <- data.frame(plot,bird,area)
birdlist
```

```
##   plot bird area
## 1    1    a   10
## 2    1    b   10
## 3    2    a    5
## 4    2    b    5
## 5    2    c    5
## 6    2    d    5
## 7    3    a   15
## 8    3    b   15
## 9    3    c   15
```

```
# summarize the following data frame to a summary table.
# option 1
birdlist %>%
  group_by(plot) %>%
  summarise(bird = n(), area = unique(area))
```

```
## # A tibble: 3 x 3
##   plot bird area
##   <dbl> <int> <dbl>
## 1     1     2   10
## 2     2     4    5
## 3     3     3   15
```

```
# option 2
birdlist %>%
  count(plot, area, name = "bird")
```

```
##   plot area bird
## 1    1   10    2
## 2    2    5    4
## 3    3   15    3
```

```
gapminder %>%
  summarise(mean(lifeExp))
```

```
## # A tibble: 1 x 1
##   `mean(lifeExp)`
##           <dbl>
## 1           59.5
```

```
gapminder %>%
  summarise(range(lifeExp))
```

```
## # A tibble: 2 x 1
##   `range(lifeExp)`
##           <dbl>
## 1           23.6
## 2           82.6
```

```
gapminder %>%
  filter(country=="India") %>%
  group_by(country) %>%
  summarise(GDPmax=max(gdpPercap),
            GDPmin=min(gdpPercap),
            GDPmean=mean(gdpPercap))
```

```
## # A tibble: 1 x 4
##   country GDPmax GDPmin GDPmean
##   <fct>    <dbl> <dbl>   <dbl>
## 1 India   2452.   547.  1057.
```

## count/summarize

```
library(tidyverse)
plot <- c(rep(1,2), rep(2,4), rep(3,3))
bird <- as.factor(c('a','b', 'a','b', 'c', 'd', 'a', 'b', 'c'))
area <- c(rep(10,2), rep(5,4), rep(15,3))

birdlist <- data.frame(plot,bird,area)
birdlist
```

```
##   plot bird area
## 1    1    a  10
## 2    1    b  10
## 3    2    a   5
## 4    2    b   5
## 5    2    c   5
## 6    2    d   5
## 7    3    a  15
## 8    3    b  15
## 9    3    c  15
```

```
#birdlist %>%   group_by(plot, area) %>%   mutate(count(bird))
```

```
birdlist %>%
  group_by(plot, area) %>%
  dplyr::summarize(bird = n(), # when summarize doesn't work directly use it (dplyr::)like this
                  .groups = "drop") # to summarize of a column with reference to two other variables.
```

```
## # A tibble: 3 x 3
##   plot area bird
```



```
##      <dbl> <dbl> <int>
## 1      1      10      2
## 2      2       5      4
## 3      3     15      3
```

## count sites

```
treatment <- c(rep('ab',2), rep('bgrnf', 8), rep('bgpnf', 4))
site <- c('ab1', 'ab2',
         rep('bgrnf1', 3),
         rep('bgrnf2', 2),
         'bgrnf3',
         'bgrnf4',
         'bgrnf5',
         rep('bgpnf1', 2),
         rep('bgpnf2', 2))
data <- data.frame(treatment, site)
library(tidyverse)
# to find the site per each treatment
data %>% group_by(treatment) %>% count(treatment)
```

```
## # A tibble: 3 x 2
## # Groups:   treatment [3]
##   treatment      n
##   <chr>      <int>
## 1 ab          2
## 2 bgpnf       4
## 3 bgrnf       8
```

## separate

text to columns

```
df <- data.frame(films = c("Spider_man", "James_bond", "Iron_man", "Bat_man"))
df
```

```
##      films
## 1 Spider_man
## 2 James_bond
## 3 Iron_man
## 4 Bat_man
```

```
df1 <- df %>%
  separate(films, c("a", "b"), sep='([_])')
df1
```

```
##      a      b
## 1 Spider man
## 2 James bond
## 3 Iron man
## 4 Bat man
```

## unite

```
df1 %>% unite("names", a:b, remove=FALSE)
```

```
##      names      a      b
## 1 Spider_man Spider  man
## 2 James_bond  James bond
## 3   Iron_man   Iron  man
## 4   Bat_man   Bat   man
```

## join

```
df2 <- data.frame(id=c(1:4), films = c("Spider_man", "James_bond", "Iron_man", "Bat_man"))
df3 <- data.frame(id=c(1:4), country= rep("us", 4))
```

```
df4 <- left_join(df2, df3, by="id")
df4
```

```
##   id      films country
## 1  1 Spider_man      us
## 2  2 James_bond      us
## 3  3   Iron_man      us
## 4  4   Bat_man      us
```

## across

for multiple variables

```
library(tidyverse)
srno <- c(1:2)
film <- c("arabica", "robust")
rate <- c("good", "better")
lang_Eng <- c("yes", "yes")

films <- data.frame(srno, film, rate, lang_Eng)

str(films)
```

```
## 'data.frame':   2 obs. of  4 variables:
## $ srno      : int  1 2
## $ film      : chr  "arabica" "robust"
## $ rate      : chr  "good" "better"
## $ lang_Eng  : chr  "yes" "yes"
```

```
films <- films %>%
  mutate(across(c(rate, lang_Eng), as.factor))

str(films)
```

```
## 'data.frame': 2 obs. of 4 variables:
## $ srno : int 1 2
## $ film : chr "arabica" "robust"
## $ rate : Factor w/ 2 levels "better","good": 2 1
## $ lang_Eng: Factor w/ 1 level "yes": 1 1
```

## everything

select a key variable and everything

```
library(gapminder)
gapminder %>% select(pop, everything())
```

```
## # A tibble: 1,704 x 6
##       pop country    continent  year lifeExp gdpPercap
##   <int> <fct>      <fct>    <int> <dbl>    <dbl>
## 1  8425333 Afghanistan Asia      1952  28.8    779.
## 2  9240934 Afghanistan Asia      1957  30.3    821.
## 3 10267083 Afghanistan Asia      1962  32.0    853.
## 4 11537966 Afghanistan Asia      1967  34.0    836.
## 5 13079460 Afghanistan Asia      1972  36.1    740.
## 6 14880372 Afghanistan Asia      1977  38.4    786.
## 7 12881816 Afghanistan Asia      1982  39.9    978.
## 8 13867957 Afghanistan Asia      1987  40.8    852.
## 9 16317921 Afghanistan Asia      1992  41.7    649.
##10 22227415 Afghanistan Asia      1997  41.8    635.
## # ... with 1,694 more rows
```

## toupper

## tolower

```
library(stringr)

data <- data.frame(Dose.Cm=c("d1", "D2", "D3"),
                  Len.km=c("High", 'low', 'Low'))
glimpse(data)
```

```
## Rows: 3
## Columns: 2
## $ Dose.Cm <chr> "d1", "D2", "D3"
## $ Len.km <chr> "High", "low", "Low"
```

```
data %>% mutate(Dose.Cm= tolower(Dose.Cm), Len.km=toupper(Len.km))
```

```
##   Dose.Cm Len.km
## 1     d1  HIGH
## 2     d2   LOW
## 3     d3   LOW
```

## factor

```
data <- data.frame(Dose.Cm=c("d1", "D2", "D3"),
                  Len.km=c("high", 'low', 'medium'))
data <- data %>% mutate(len= as.factor(Len.km))

glimpse(data)
```

```
## Rows: 3
## Columns: 3
## $ Dose.Cm <chr> "d1", "D2", "D3"
## $ Len.km <chr> "high", "low", "medium"
## $ len <fct> high, low, medium
```

## change order of factor

```
data %>% mutate(len= fct_relevel(len, c('low', 'medium', 'high')))
```

```
##   Dose.Cm Len.km   len
## 1      d1  high  high
## 2      D2   low   low
## 3      D3 medium medium
```

## parse\_number

This drops any non-numeric characters before or after the first number. The grouping mark specified by the locale is ignored inside the number.

```
library(tidyverse)
class <- c('8th', '9th', '10th')
students <- c('25-30', '35-41', '21-28')
school <- data.frame(class, students)
school
```

```
##   class students
## 1   8th    25-30
## 2   9th    35-41
## 3  10th    21-28
```

```
glimpse(school) # notice students is a binned variable it is not a numeric.
```

```
## Rows: 3
## Columns: 2
## $ class <chr> "8th", "9th", "10th"
## $ students <chr> "25-30", "35-41", "21-28"
```

```
school %>% mutate(students= parse_number(students)) %>% glimpse()
```

```
## Rows: 3  
## Columns: 2  
## $ class    <chr> "8th", "9th", "10th"  
## $ students <dbl> 25, 35, 21
```

```
school %>% mutate(students= parse_number(students))
```

```
##   class students  
## 1   8th       25  
## 2   9th       35  
## 3  10th       21
```

```
# now students because number with first value of the column
```

## pivot longer

```
library(tidyverse)
```

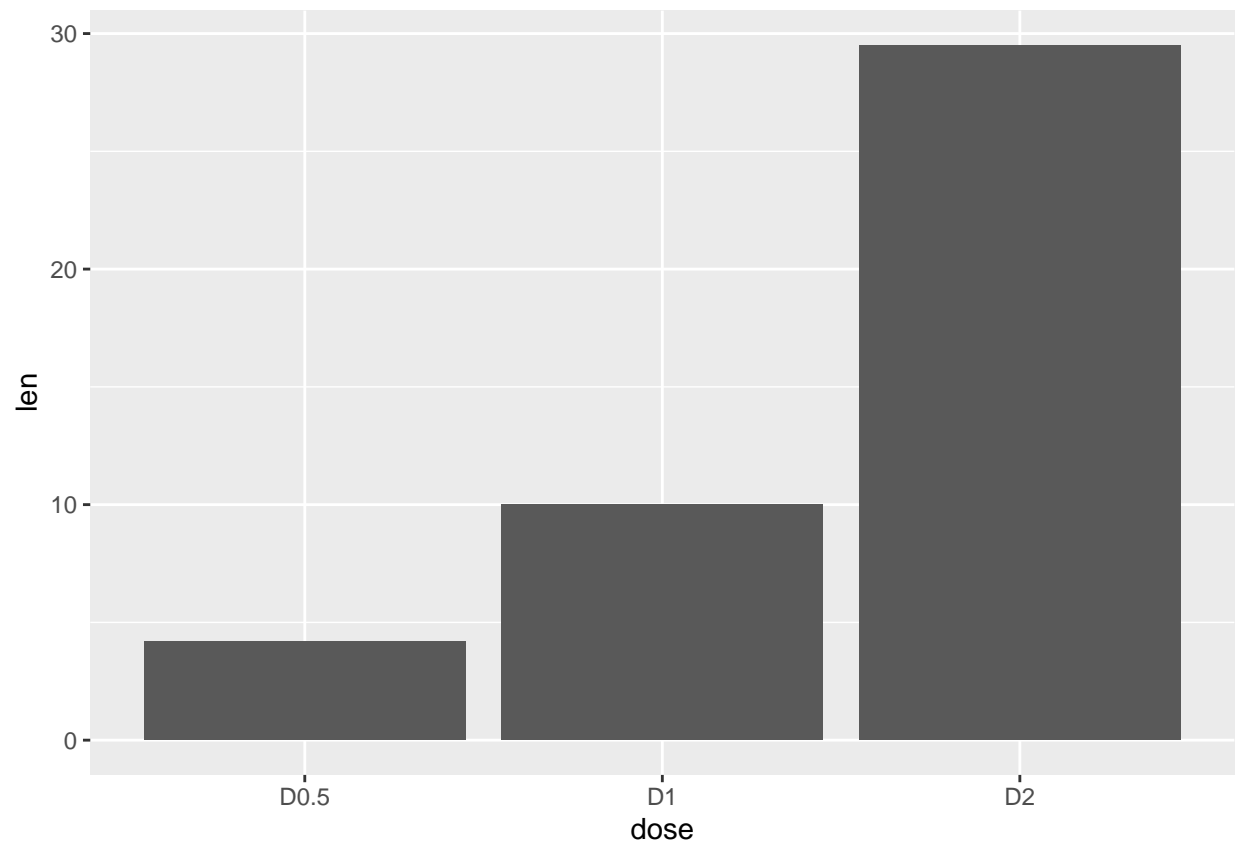
```
rawdata <- data.frame(species_1=rnorm(n = 40, mean = 300, sd = 18.5), species_2=rnorm(40, 305, 16.7))
```

```
data <- pivot_longer(data = rawdata, cols = species_1:species_2, names_to = 'species', values_to = 'weight')
```

## ggplot

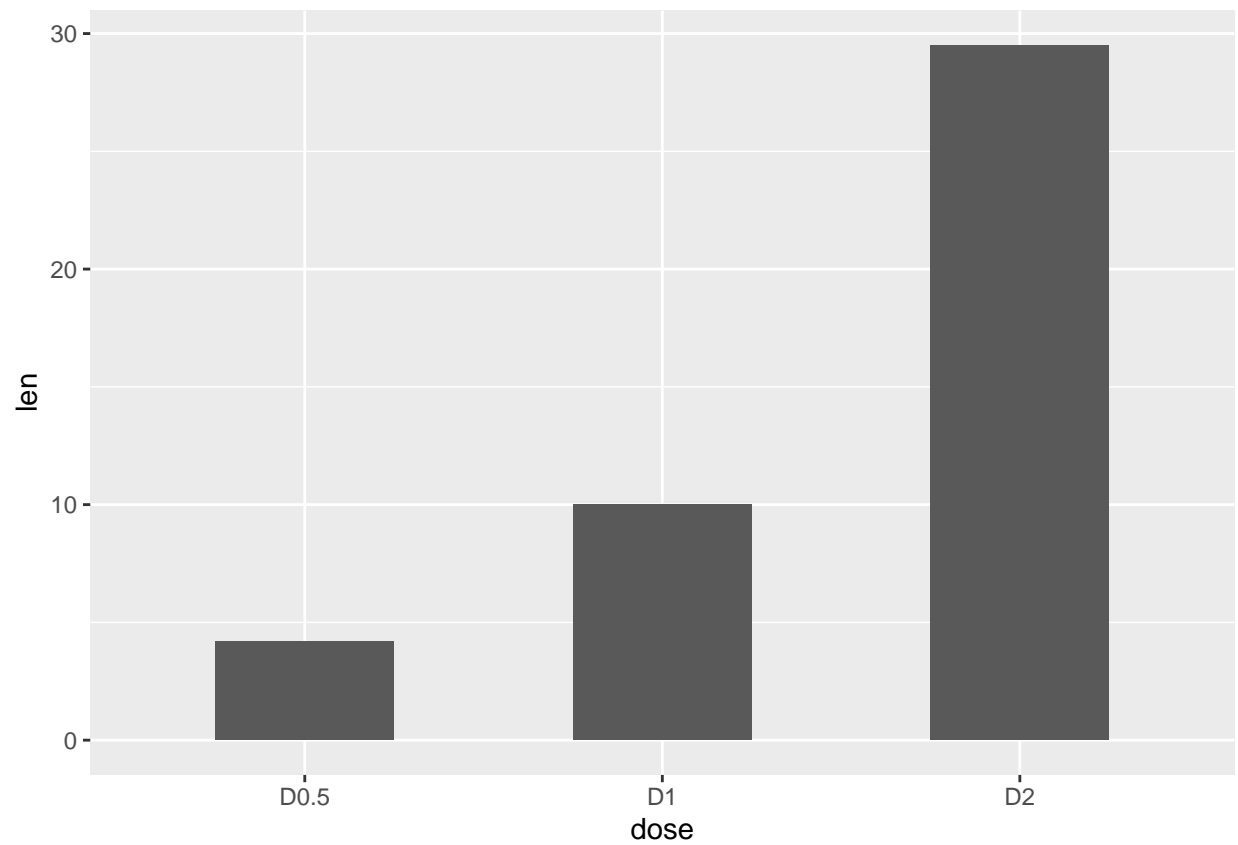
```
#sthda.com/english/wiki/ggplot2-barplots-quick-start-guide-r-software-and-data-visualization  
df <- data.frame(dose=c("D0.5", "D1", "D2"),  
                 len=c(4.2, 10, 29.5))
```

```
library(ggplot2)  
# Basic barplot  
p<-ggplot(data=df, aes(x=dose, y=len)) +  
  geom_bar(stat="identity")  
p
```

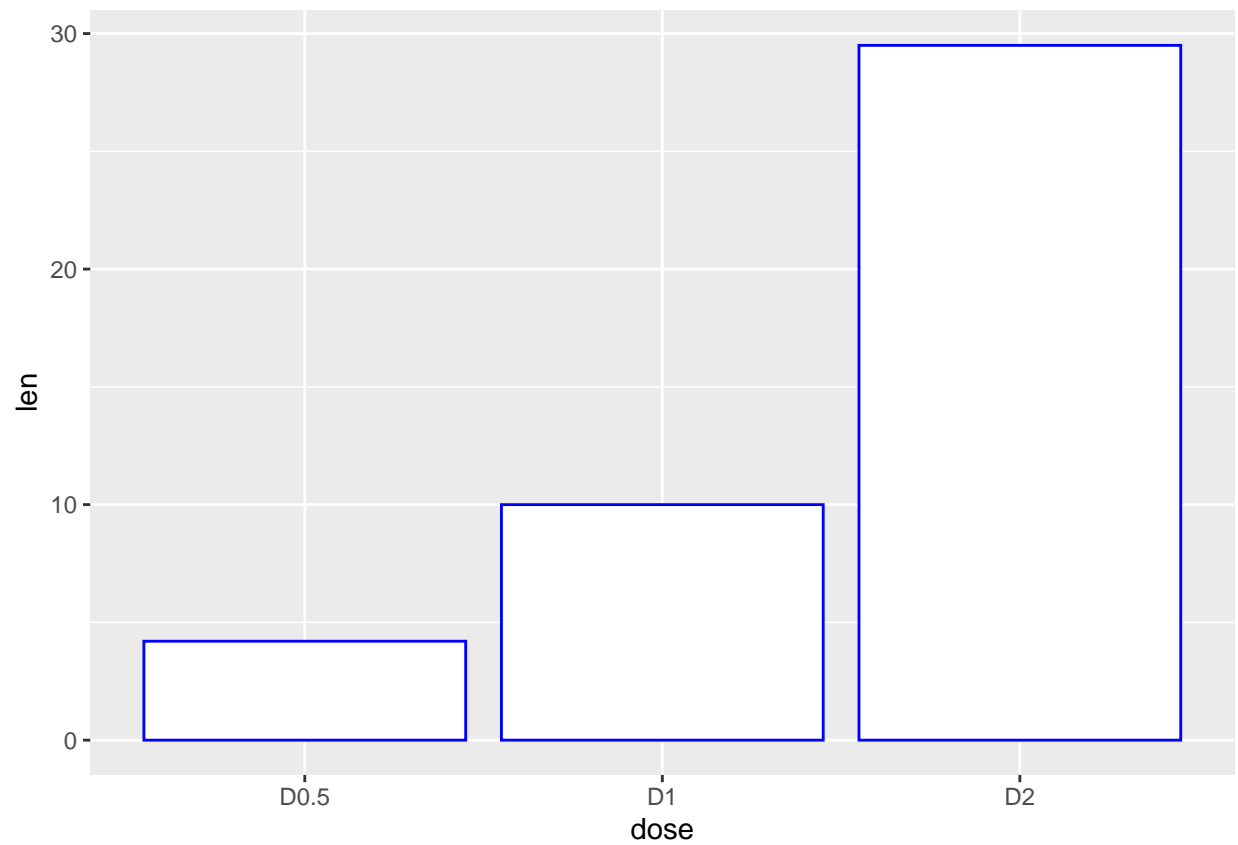


```
# Horizontal bar plot  
# p + coord_flip()
```

```
# Change the width of bars  
ggplot(data=df, aes(x=dose, y=len)) +  
  geom_bar(stat="identity", width=0.5)
```

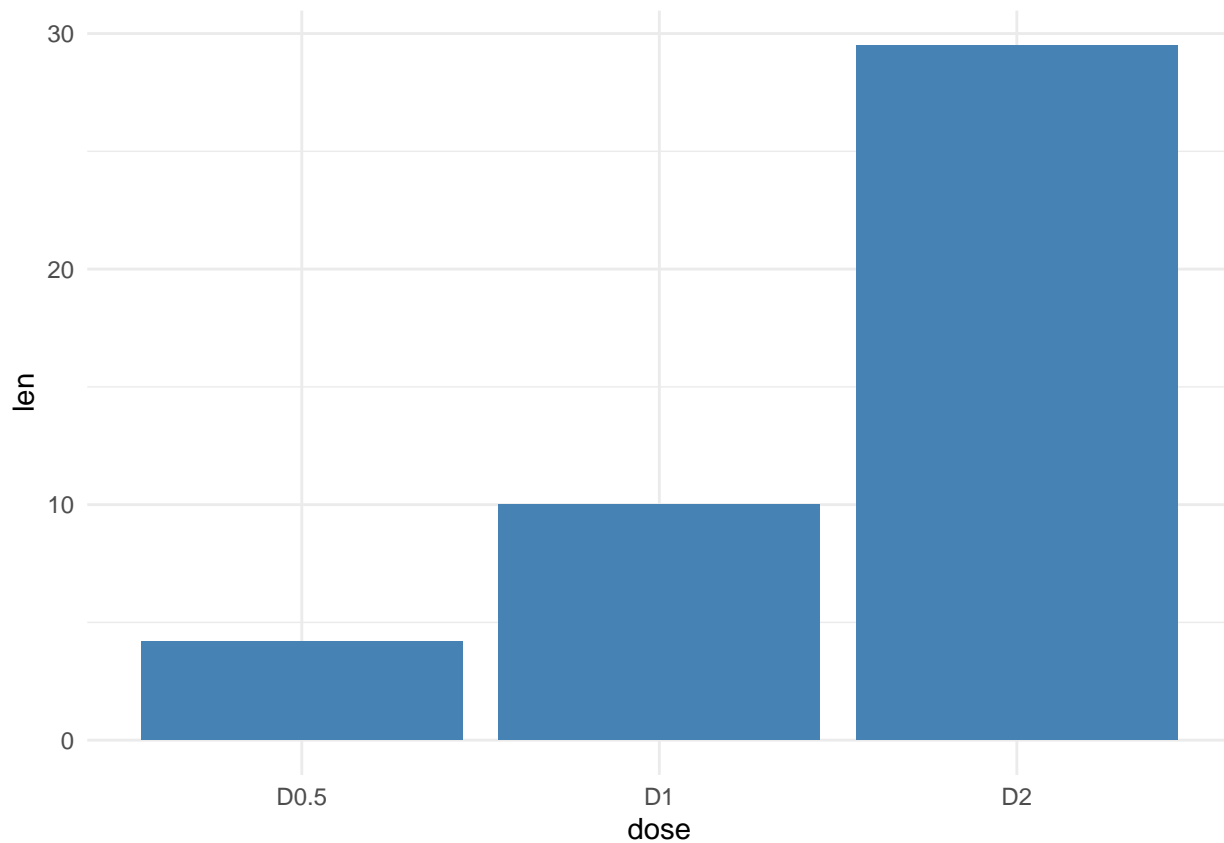


```
# Change colors  
ggplot(data=df, aes(x=dose, y=len)) +  
  geom_bar(stat="identity", color="blue", fill="white")
```



```
# Minimal theme + blue fill color
p<-ggplot(data=df, aes(x=dose, y=len)) +
  geom_bar(stat="identity", fill="steelblue")+
  theme_minimal()
p
```





## months

```
library(lubridate)
```

```
##
```

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

```
## The following objects are masked from 'package:base':
```

```
##
```

```
## date, intersect, setdiff, union
```

```
months <- seq(month(1:12)) # make months
```

```
months <- month.abb[months] # make abbreviations
```

```
temperature <- c(10,12,22,32,35,30,33,28,29,25,19,14)
```

```
myframe <- data.frame(months,temperature) # creating a new data frame
```

```
library(tidyverse)
```

```
glimpse(myframe)
```

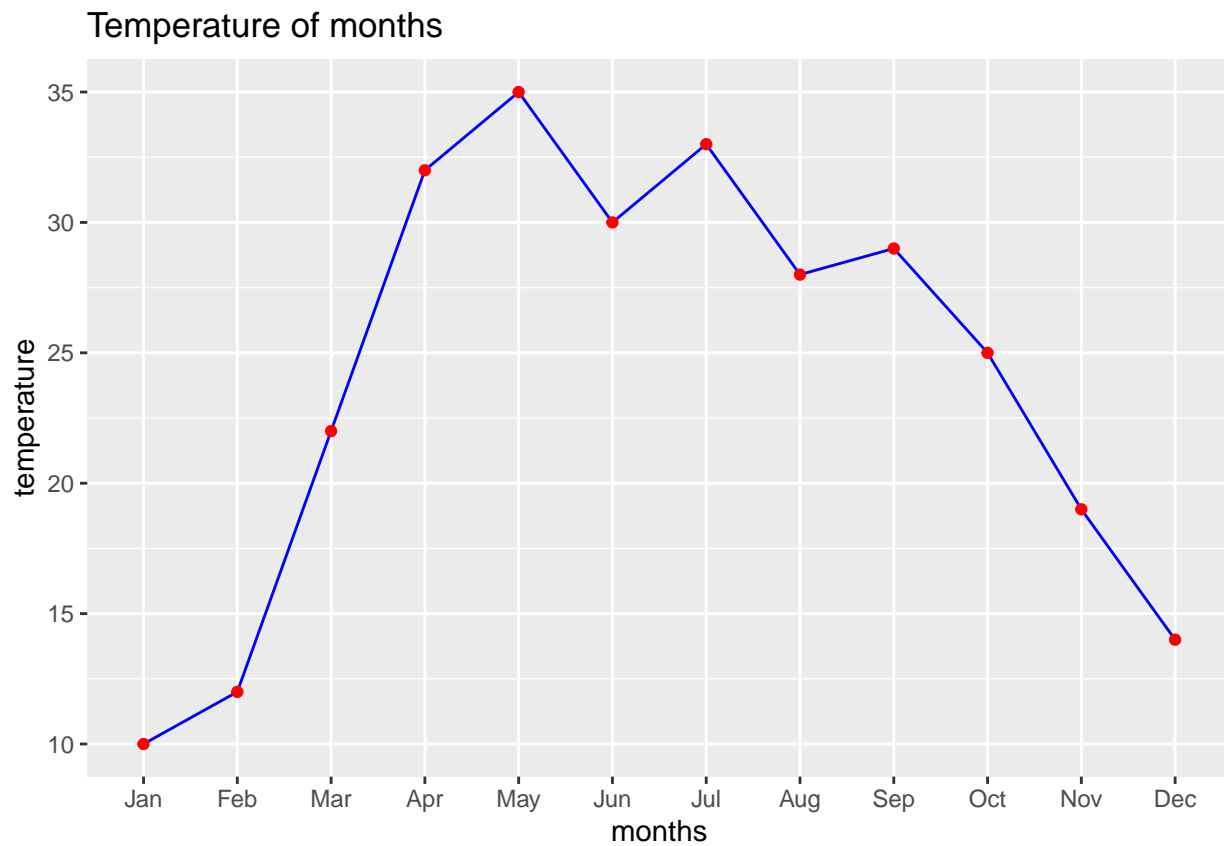
```
## Rows: 12
```

```
## Columns: 2
```

```
## $ months      <chr> "Jan", "Feb", "Mar", "Apr", "May", "Jun", "Jul", "Aug", "S~
```

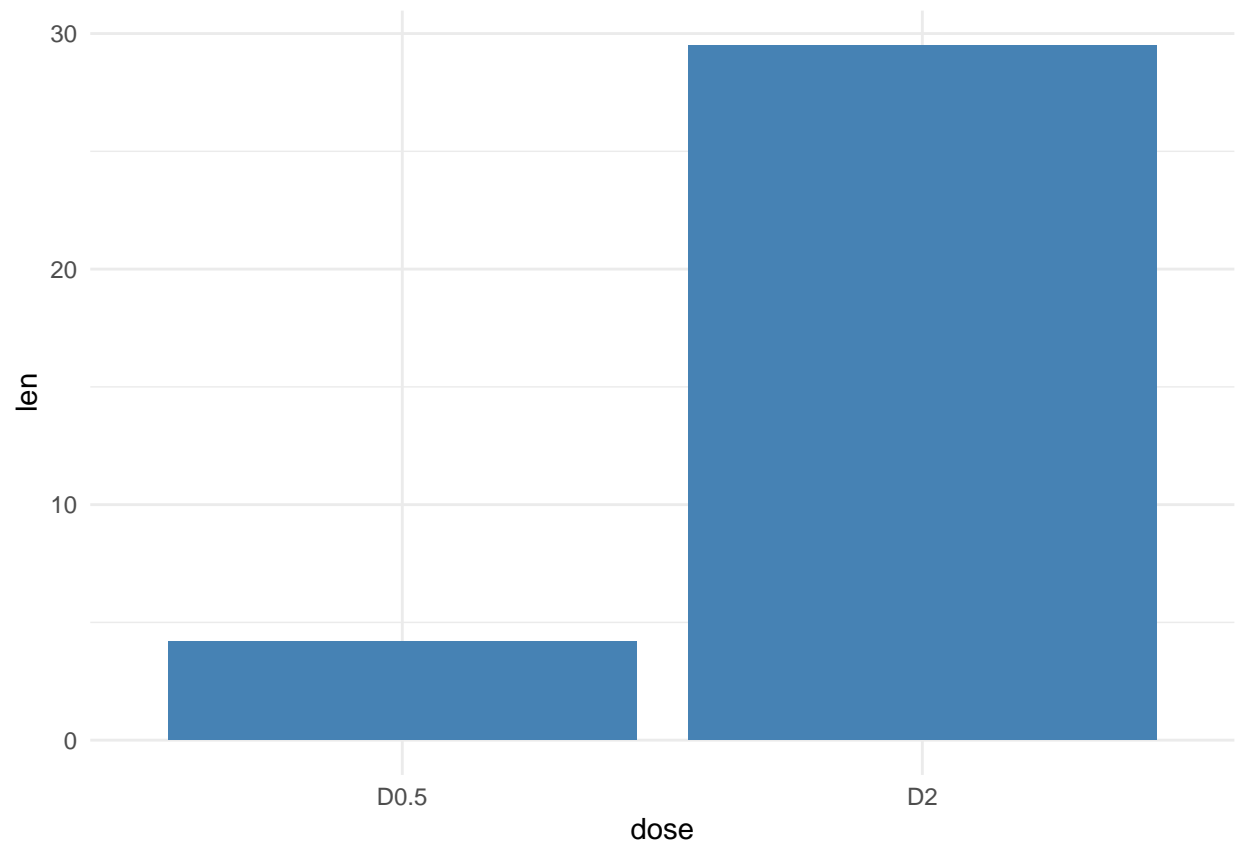
```
## $ temperature <dbl> 10, 12, 22, 32, 35, 30, 33, 28, 29, 25, 19, 14
```

```
library(ggplot2)
ggplot(myframe, aes(x=months, y=temperature, group=1))+
  geom_line(col='blue')+
  geom_point(col='red')+
  ggtitle('Temperature of months')+
  scale_x_discrete(limits = month.abb) # this will order months on the x axis
```

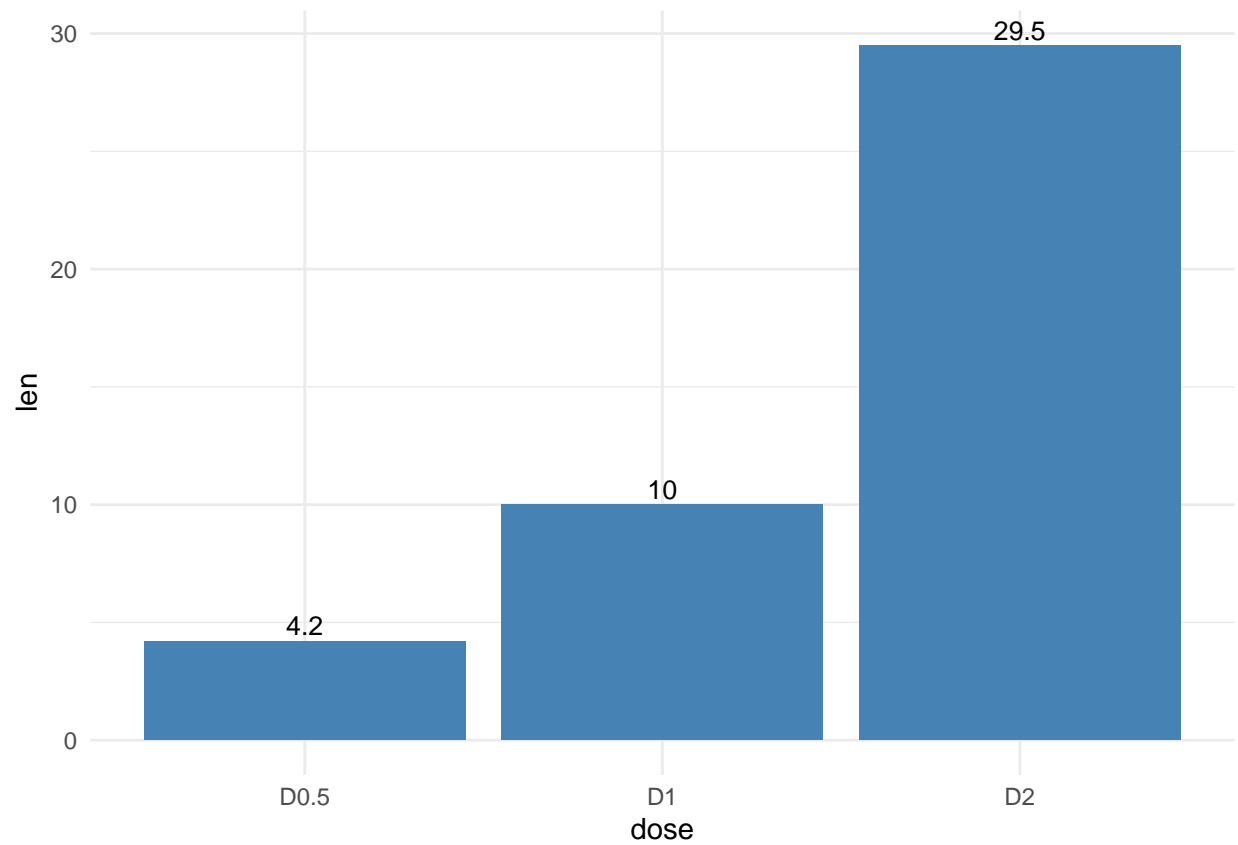


```
p + scale_x_discrete(limits=c("D0.5", "D2"))
```

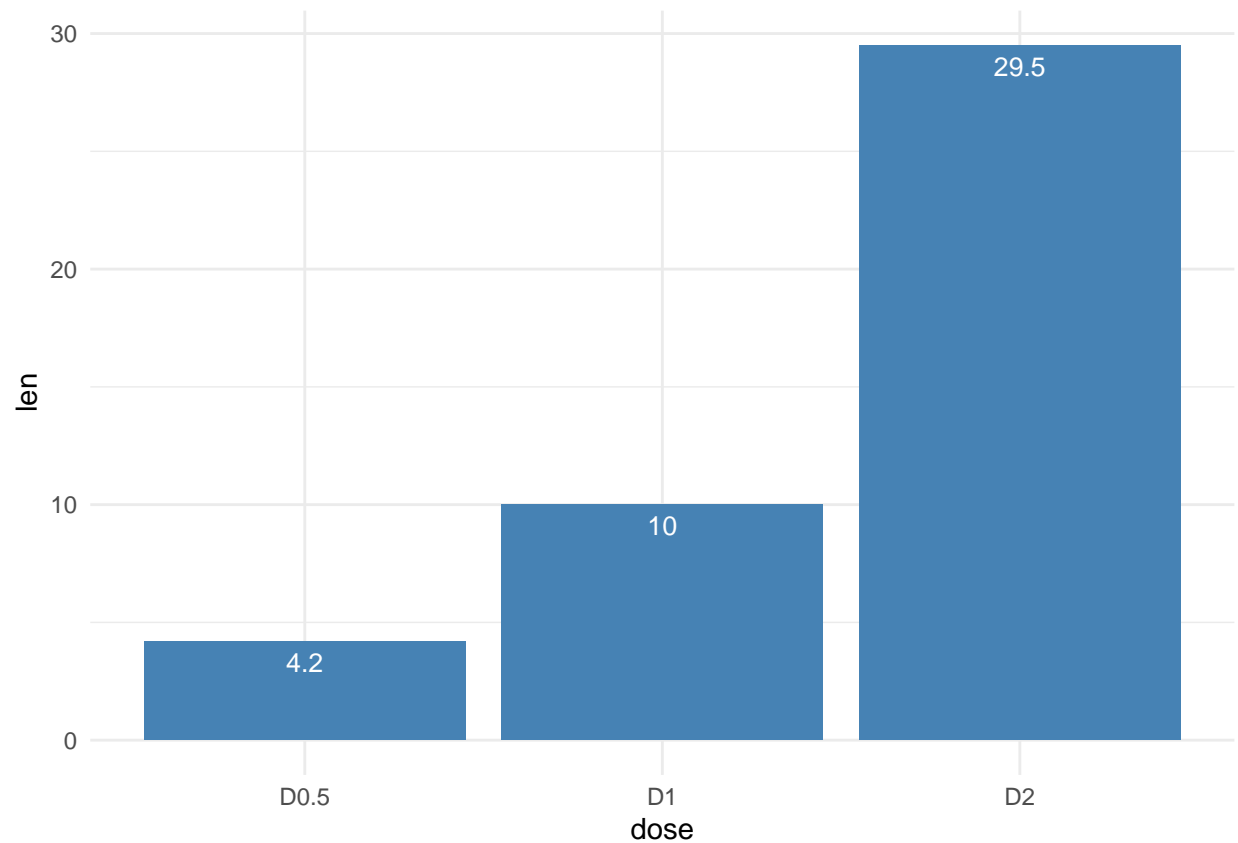
```
## Warning: Removed 1 rows containing missing values (position_stack).
```



```
# Outside bars  
ggplot(data=df, aes(x=dose, y=len)) +  
  geom_bar(stat="identity", fill="steelblue")+  
  geom_text(aes(label=len), vjust=-0.3, size=3.5)+  
  theme_minimal()
```

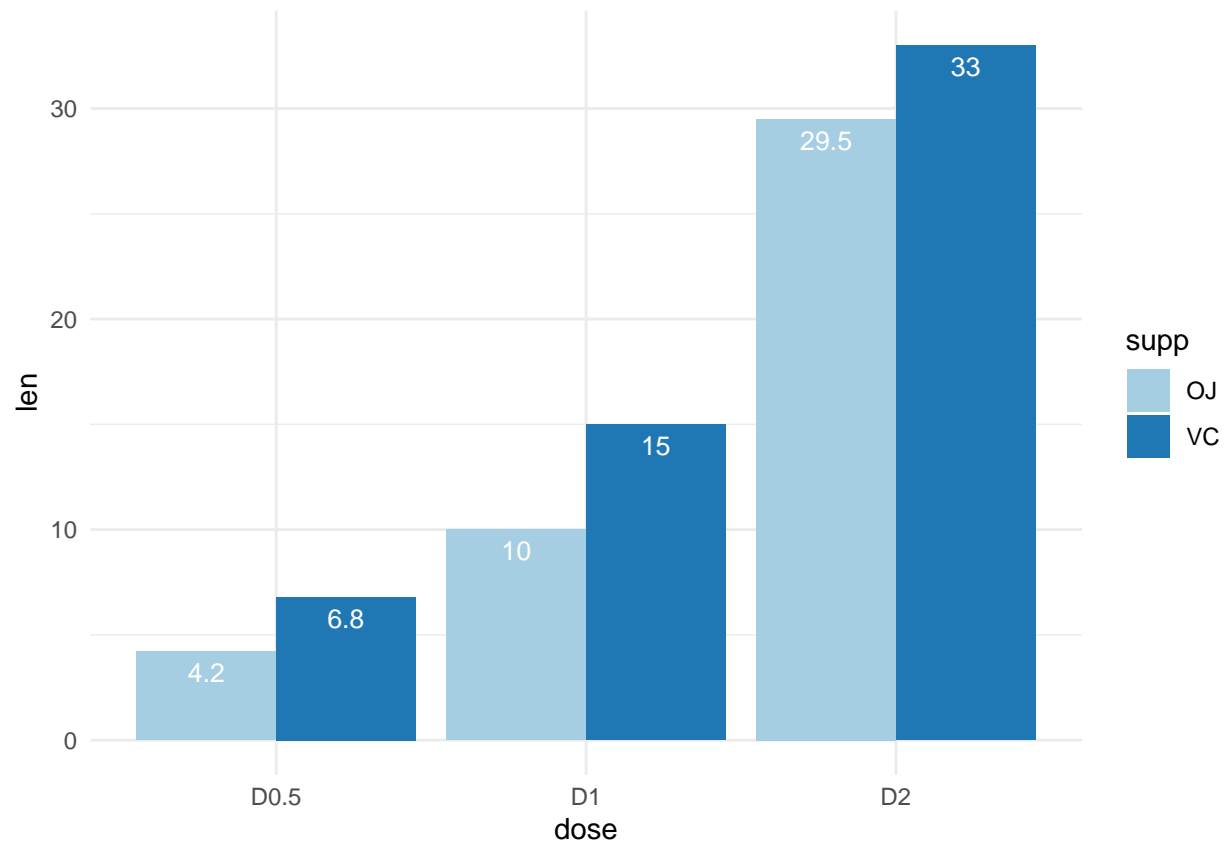


```
# Inside bars
ggplot(data=df, aes(x=dose, y=len)) +
  geom_bar(stat="identity", fill="steelblue")+
  geom_text(aes(label=len), vjust=1.6, color="white", size=3.5)+
  theme_minimal()
```

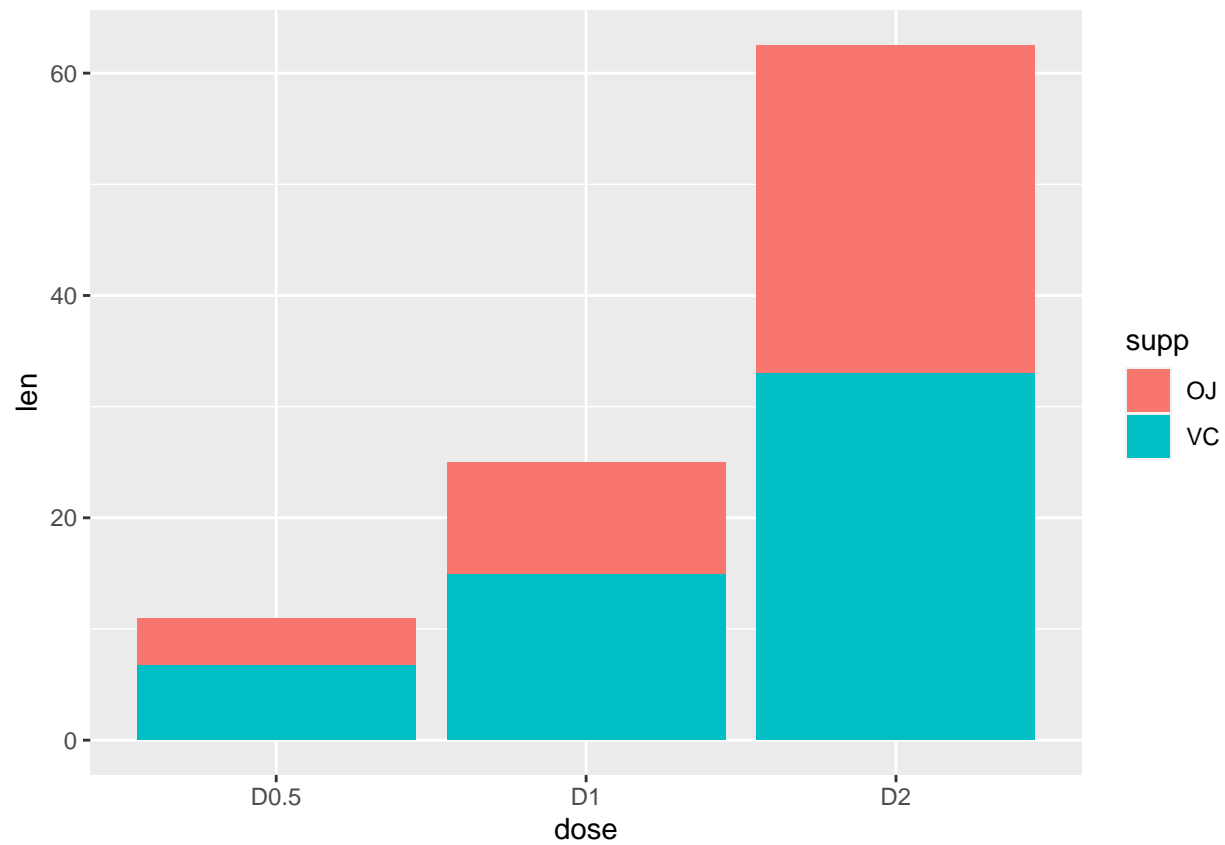


```
df2 <- data.frame(supp=rep(c("VC", "OJ"), each=3),  
                  dose=rep(c("D0.5", "D1", "D2"),2),  
                  len=c(6.8, 15, 33, 4.2, 10, 29.5))
```

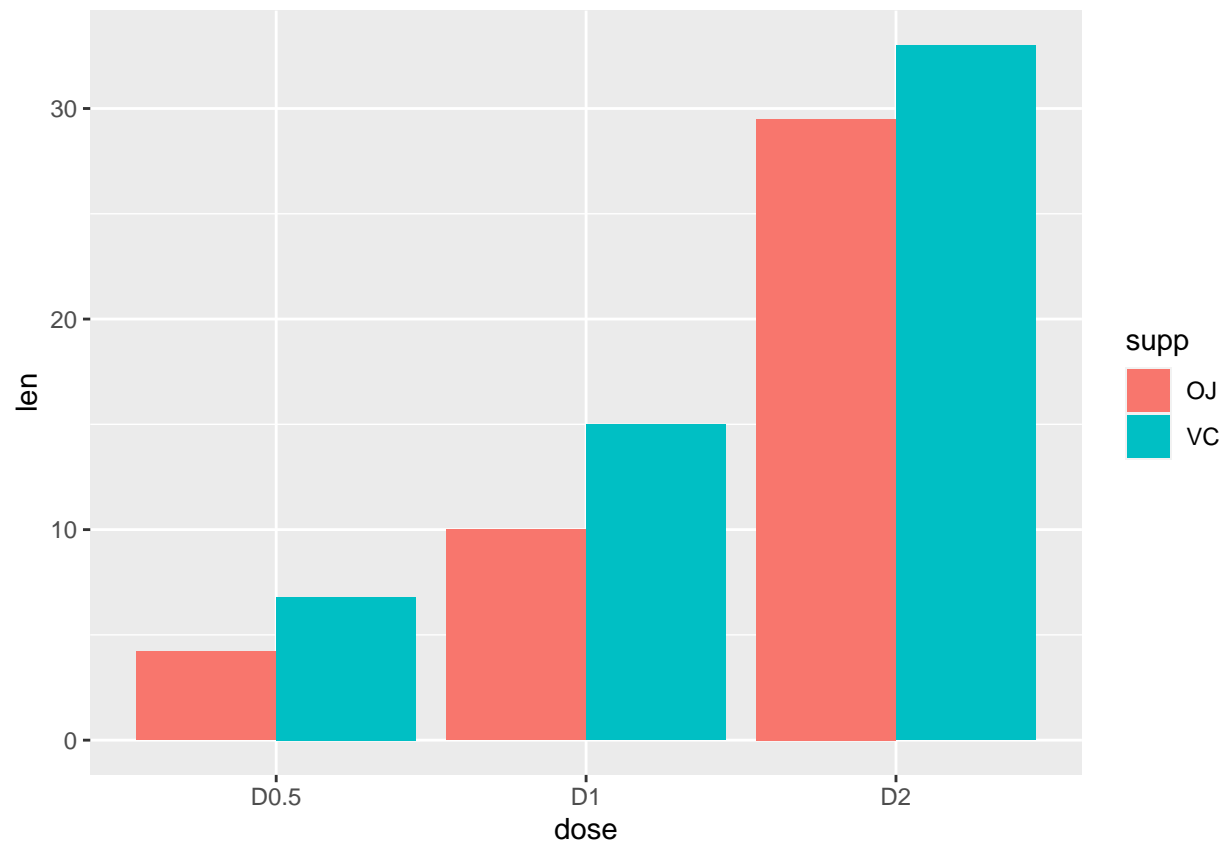
```
ggplot(data=df2, aes(x=dose, y=len, fill=supp)) +  
  geom_bar(stat="identity", position=position_dodge()) +  
  geom_text(aes(label=len), vjust=1.6, color="white",  
            position = position_dodge(0.9), size=3.5) +  
  scale_fill_brewer(palette="Paired") +  
  theme_minimal()
```



```
# Stacked barplot with multiple groups  
ggplot(data=df2, aes(x=dose, y=len, fill=supp)) +  
  geom_bar(stat="identity")
```

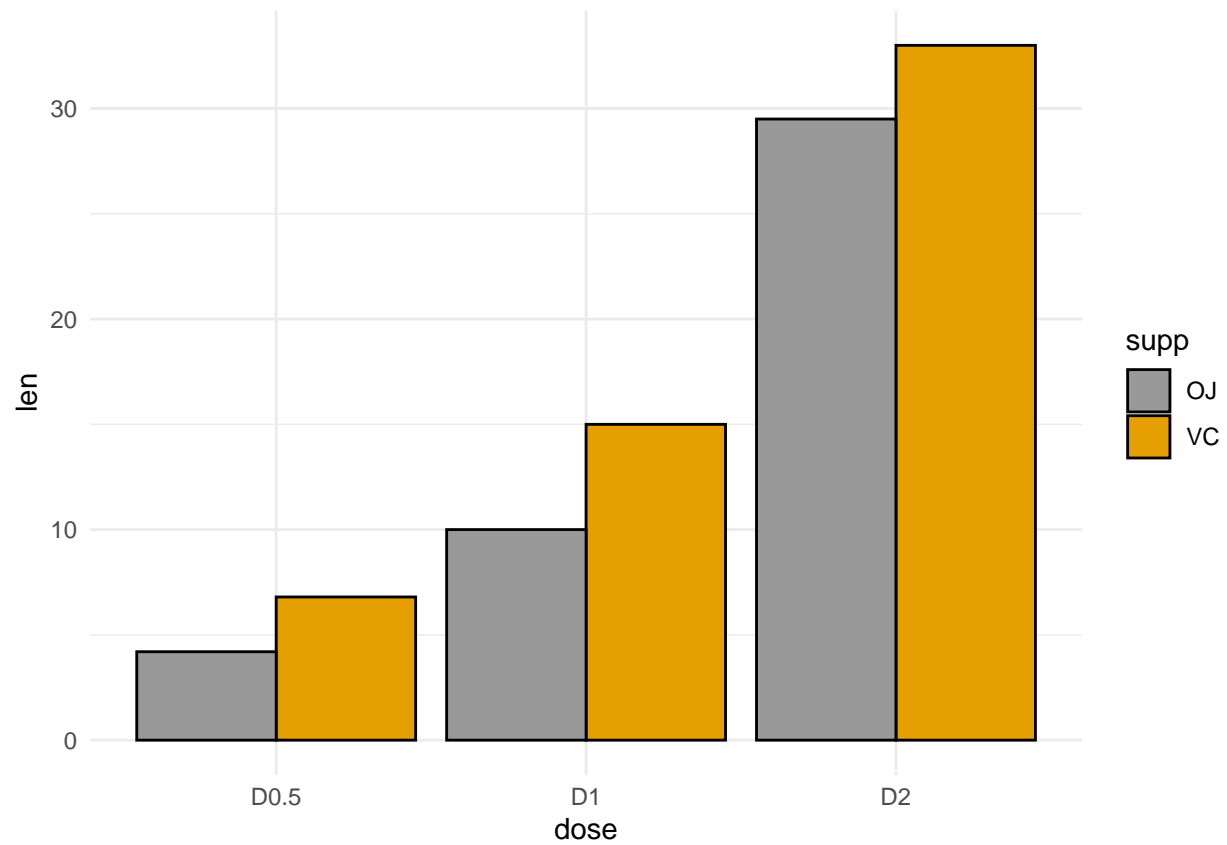


```
# Use position=position_dodge()  
ggplot(data=df2, aes(x=dose, y=len, fill=supp)) +  
geom_bar(stat="identity", position=position_dodge())
```

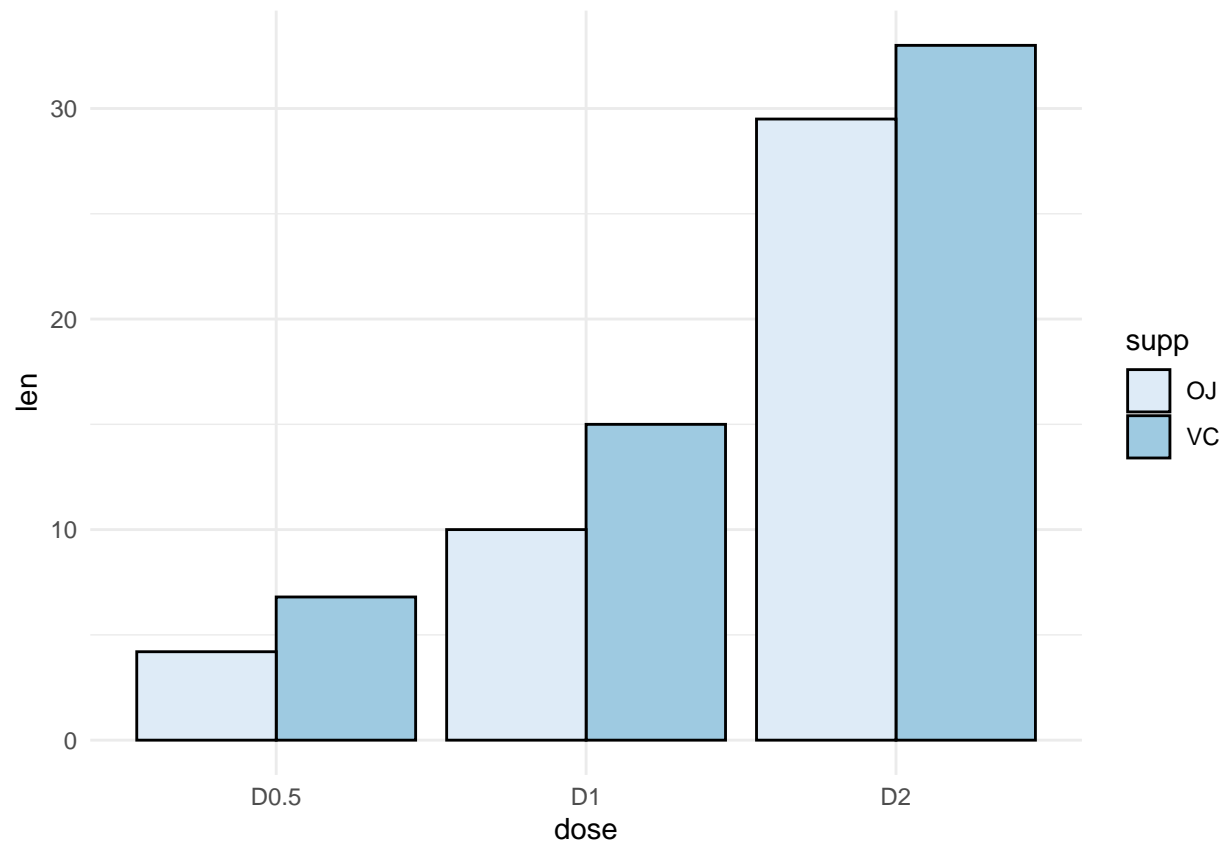


```
# Change the colors manually
p <- ggplot(data=df2, aes(x=dose, y=len, fill=supp)) +
  geom_bar(stat="identity", color="black", position=position_dodge()) +
  theme_minimal()
# Use custom colors
p + scale_fill_manual(values=c('#999999', '#E69F00'))
```





```
# Use brewer color palettes  
p + scale_fill_brewer(palette="Blues")
```



## hex plot

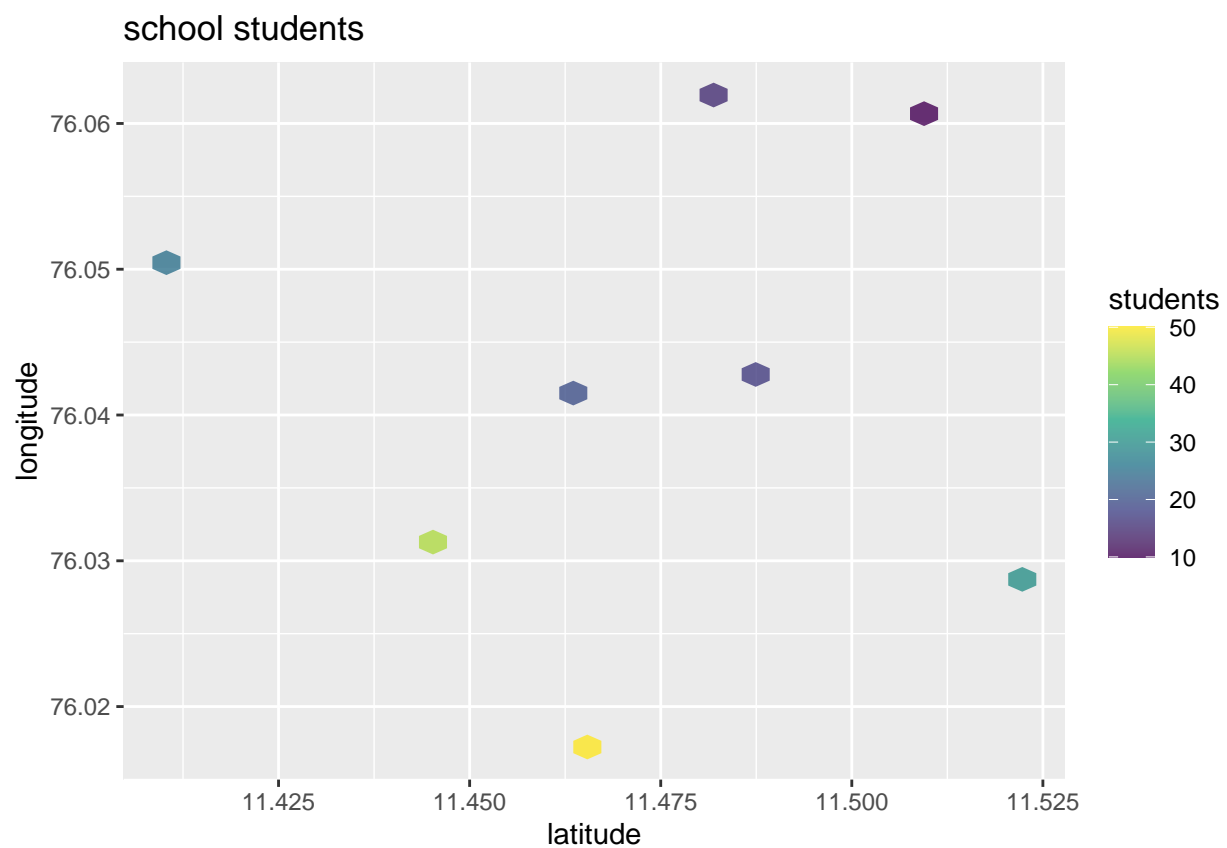
```
library(tidyverse)
# install.packages("hexbin")
class <- c(rep('10th', 8))
students <- c('10 to 15',
              '15-20',
              '17 to 24',
              '20 to 25',
              '25 to 30',
              '30 to 40',
              '45 to 47',
              '50 to 55')
latitude <- c(11.50897246,
             11.48323136,
             11.48719031,
             11.46366611,
             11.41097322,
             11.52111154,
             11.44491386,
             11.46569568)
longitude <- c(76.06032062,
              76.06192685,
```

```

76.04266851,
76.04156575,
76.05075092,
76.02846331,
76.03084141,
76.01766216)
school <- data.frame(class, students, latitude, longitude)

school %>% mutate(students= parse_number(students)) %>%
  ggplot(aes(latitude, longitude, z= students))+
  stat_summary_hex()+
  scale_fill_viridis_c(alpha= 0.8)+
  labs(fill='students', title = 'school students')

```



## Color Palettes

### Resources

- <https://colorhunt.co/>
- <https://colors.co/>
- <https://colorpalettes.net/>
- <https://www.canva.com/colors/color-palettes/>
- <https://color.adobe.com/de/create/color-wheel>
- <https://mycolor.space/>

- <http://colormind.io/>

```
#install.packages(c("tidyverse", "gapminder", "MetBrewer"))
```

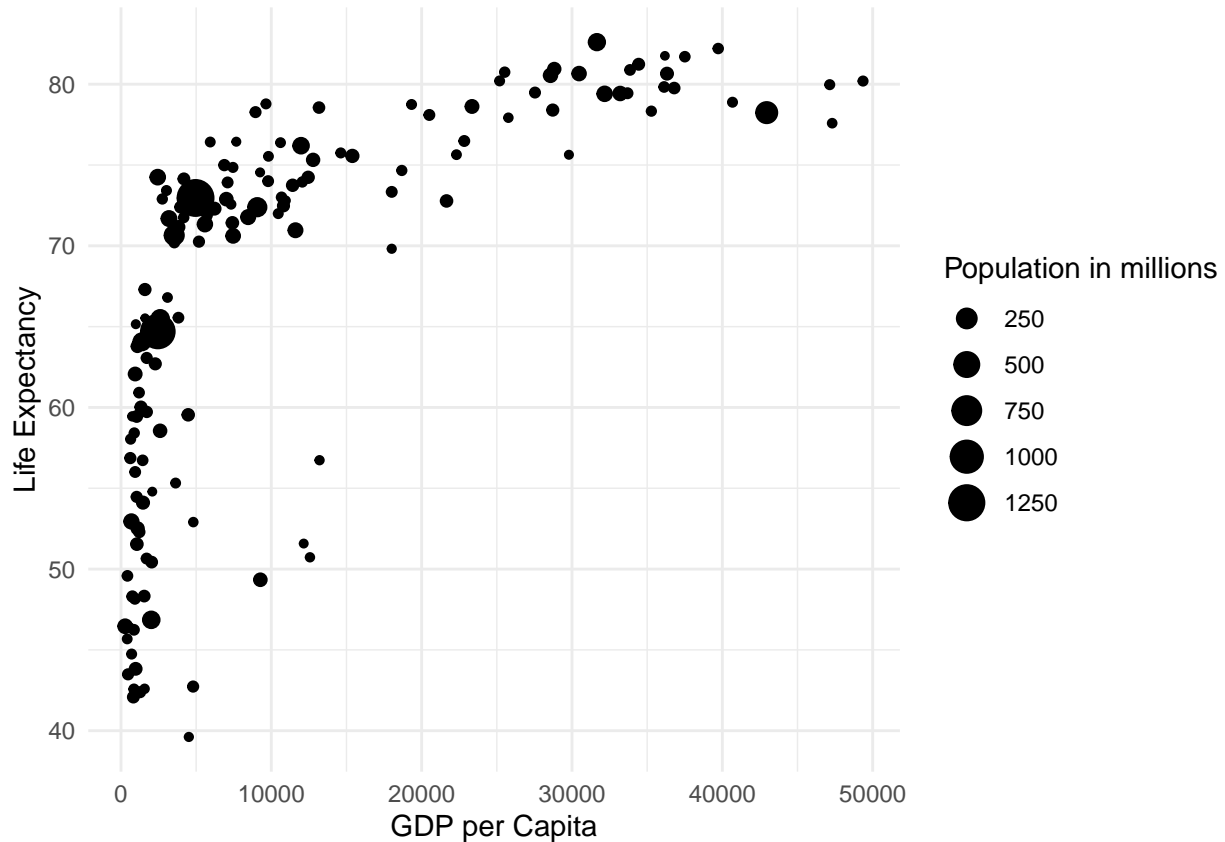
libraries

```
library(tidyverse)
library(gapminder)
# install.packages('MetBrewer')
library(MetBrewer)
```

Plot the point plot using GDP per Capita as the x- axis and LE as the y axis. Numerical variable Population to control the size of each point.

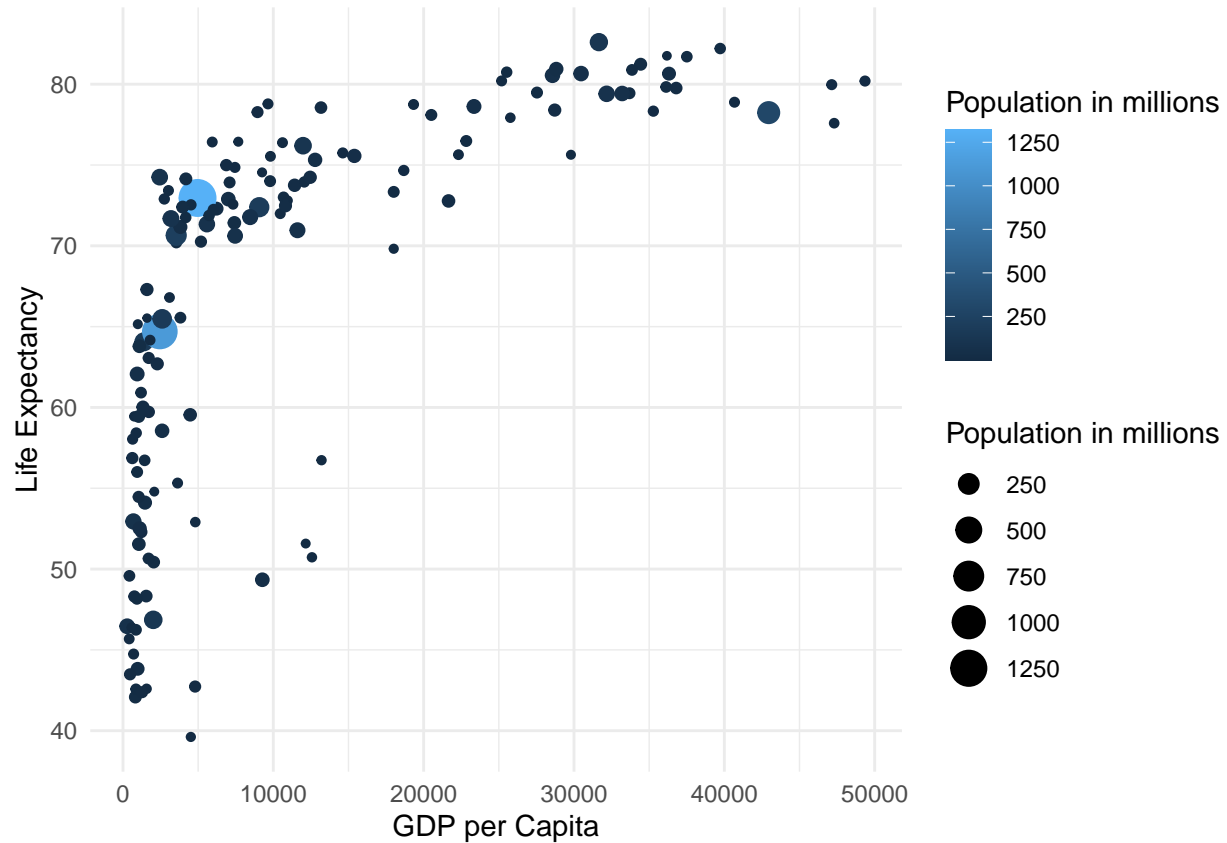
```
plot <- gapminder %>%
  filter (year==2007) %>%
  ggplot()+
  labs(x= 'GDP per Capita',
       y= 'Life Expectancy',
       color= 'Population in millions',
       size='Population in millions')+
  theme_minimal()

plot+ geom_point(aes(gdpPercap, lifeExp, size= pop/1000000))
```



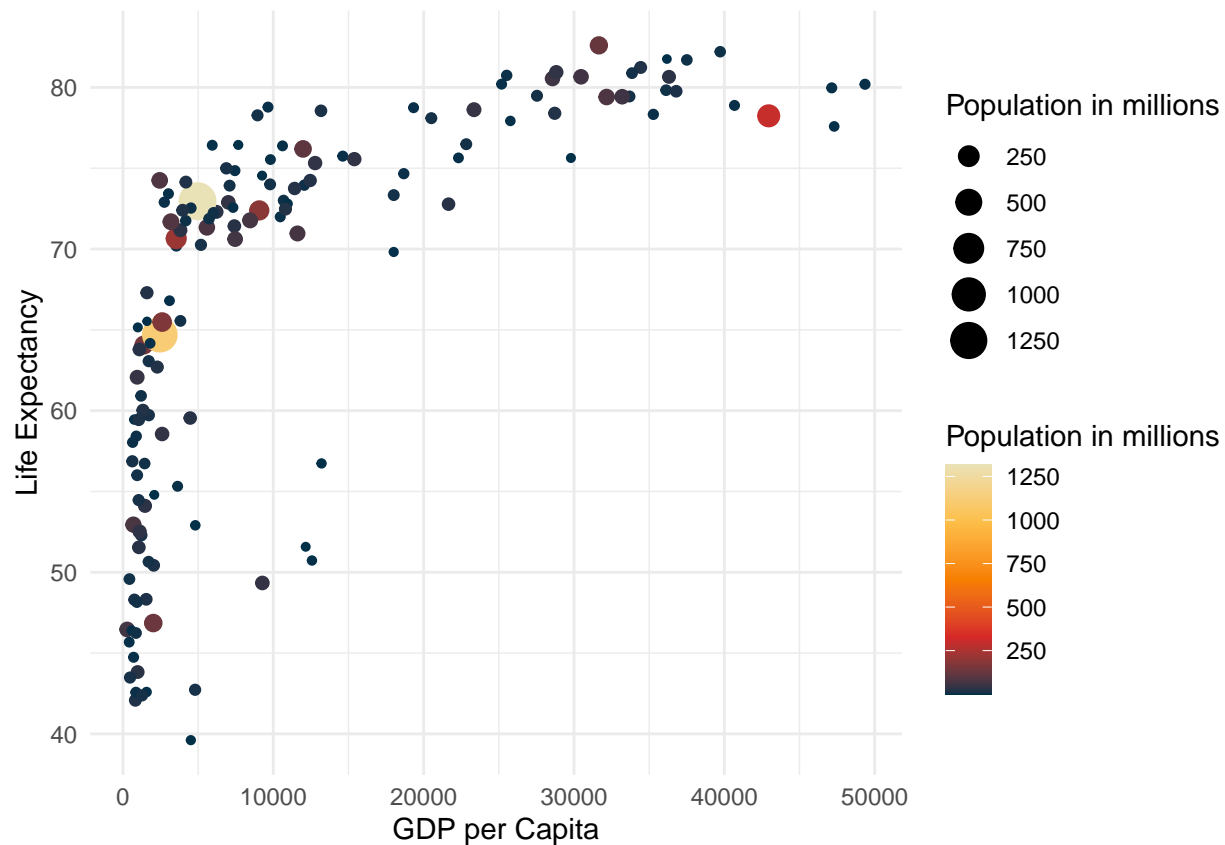
To use color in the plot, assign the Population variable to the color aesthetic. Since nothing is specified, ggplot2 chooses a color spectrum for this numerical variable (shades of blue).

```
plot + geom_point(aes(gdpPerCap, lifeExp, size= pop/1000000, color= pop/1000000))
```



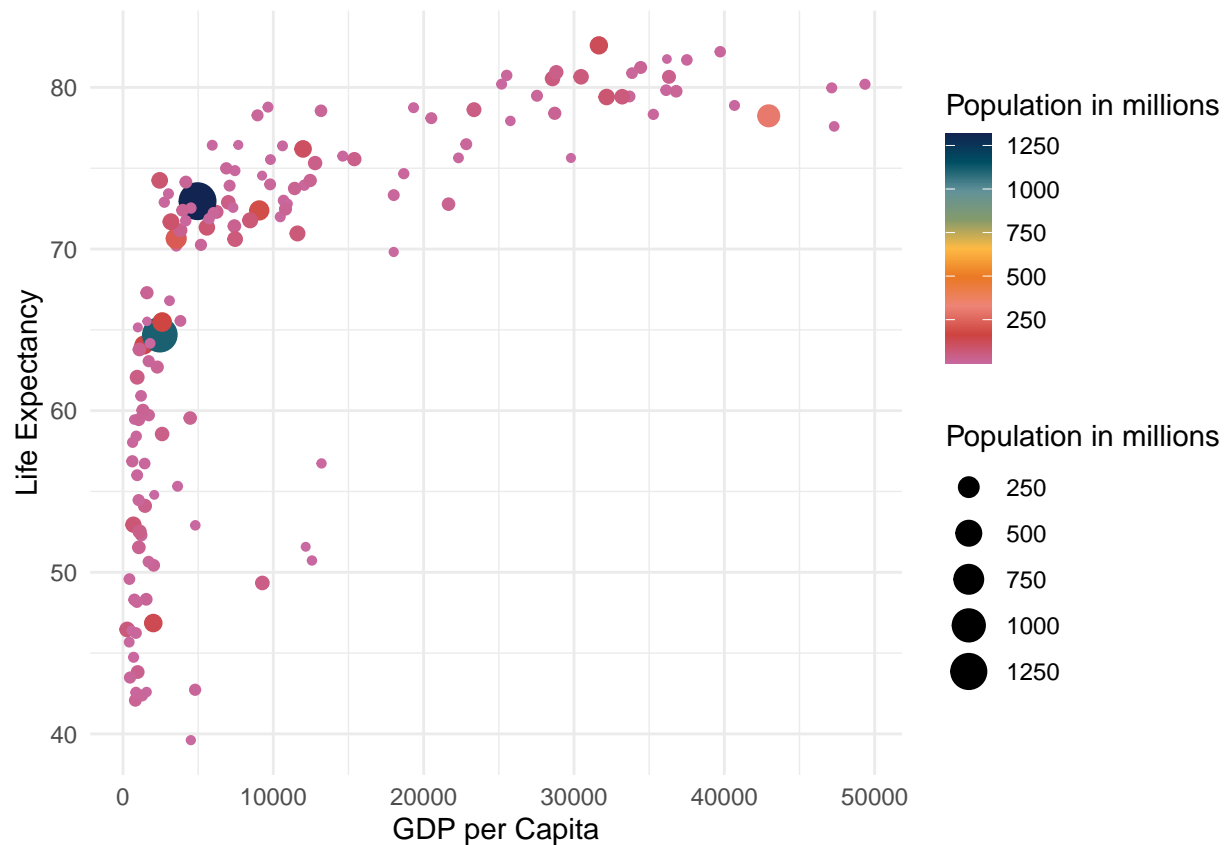
To control the color spectrum, we need to introduce a color scale. In the following plot, we have to provide a vector of hex color values. You would choose this if you got your colors from one of the mentioned above websites.

```
plot + geom_point(aes(gdpPerCap, lifeExp, size= pop/1000000, color= pop/1000000))+
  scale_color_gradientn(colors = c("#003049", "#D62828", "#F77F00", "#FCBF49", "#EAE2B7"))
```



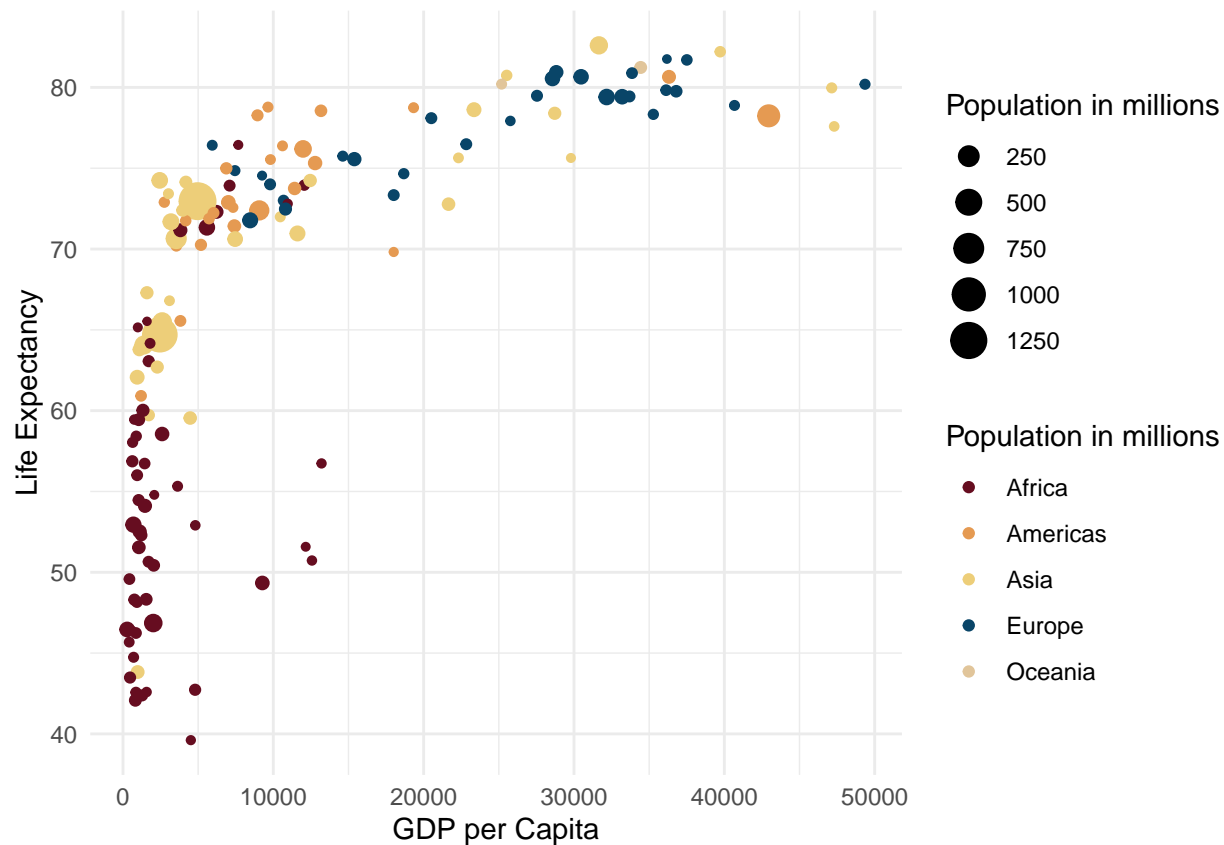
To apply one of the MetBrewer palettes, replace the hex-vector with a MetBrewer function. Within the function call, you provide the palette's name, then several colors, and tell it that we need a continuous palette since it is a numerical variable.

```
plot + geom_point(aes(gdpPercap, lifeExp, size= pop/1000000, color= pop/1000000))+
  scale_color_gradientn(colors = met.brewer('Cross', n=500, type = 'continuous'))
```



You might also want to use color palettes with non-numerical variables. Let us assume we want to apply color to the Continent variable. This implies using a manual color scale and providing a MetBrewer palette.

```
plot + geom_point(aes(gdpPercap, lifeExp, size= pop/1000000, color= continent))+
  scale_color_manual(values = met.brewer('Navajo', 5))
```



Please note if you want to apply color to the fill aesthetic rather than the color aesthetic, consider using the `scale_fill_manual` function instead of the `scale_color_manual`. This is useful for boxplots or bar charts.

```
box <- gapminder %>%
  filter(gdpPercap < 60000) %>%
  ggplot(aes(continent, gdpPercap, color= year, fill= continent))+
    geom_boxplot()+
    theme_minimal()+ labs( x= 'Continent', y= 'GDP per Capita', fill= 'Continent')
```

## scale fill manual

## themes

```
df <- data.frame(
  Names=as.factor(c('Bacteria', 'Yeast', 'None')),
  Quantity=c(2.5, 5.5, 7.5))

library(ggplot2)
library(tidyverse)
df <- df %>% mutate(Names= fct_relevel(Names, c('Bacteria', 'Yeast', 'None')))

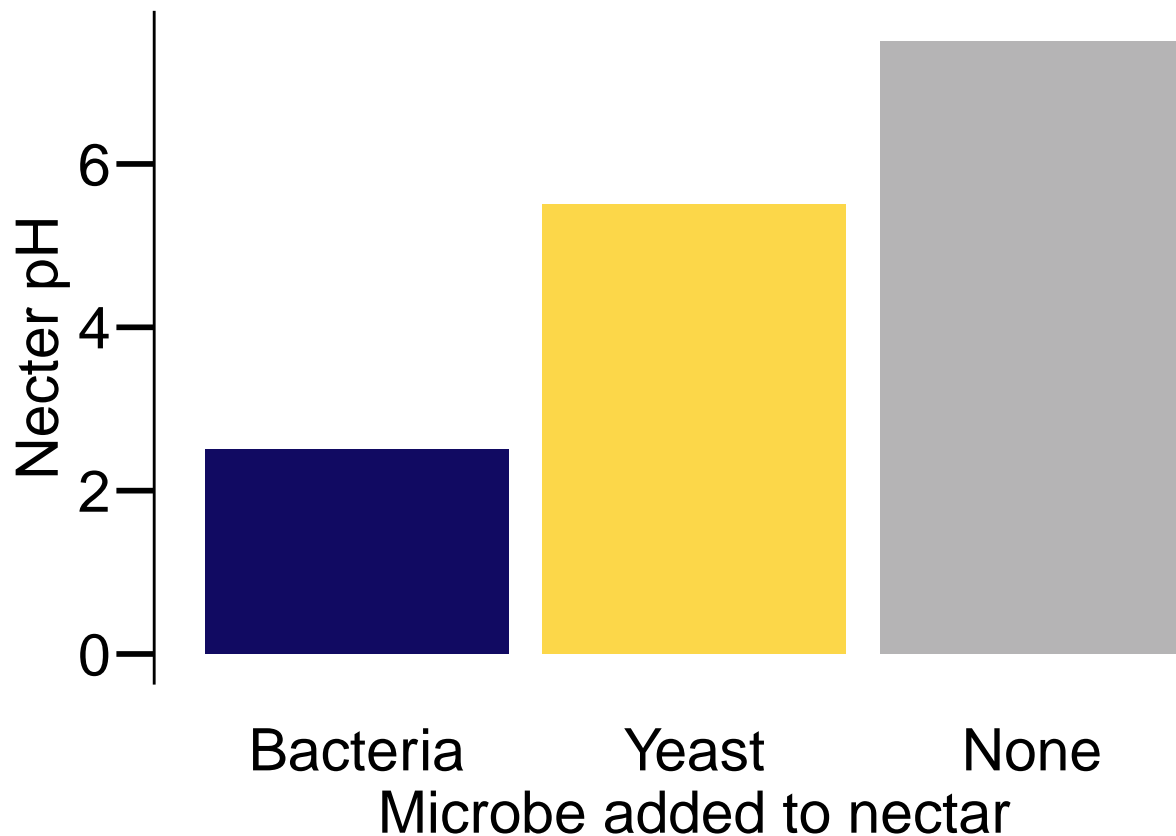
ggplot(df, aes(Names, Quantity, fill= Names))+
```



```

geom_bar(stat = 'identity')+
  scale_fill_manual(values = c('#110a62', '#fcd749', '#b5b4b5'))+
  labs(y='Nectar pH', x= 'Microbe added to nectar')+
  theme_classic()+
  theme(legend.position = 'none', axis.ticks.x = element_blank()+
  theme(axis.text = element_text(size = 22, color= 'black'))+
  theme(axis.line.x = element_blank()+
  theme(axis.ticks = element_line(size = 1, color="black"),
  axis.ticks.length = unit(.5, "cm"))+
  theme(text = element_text(size = 22))

```



graphics

```

x11() # opne a new window for graphics
graphics.off() # close the new window

```

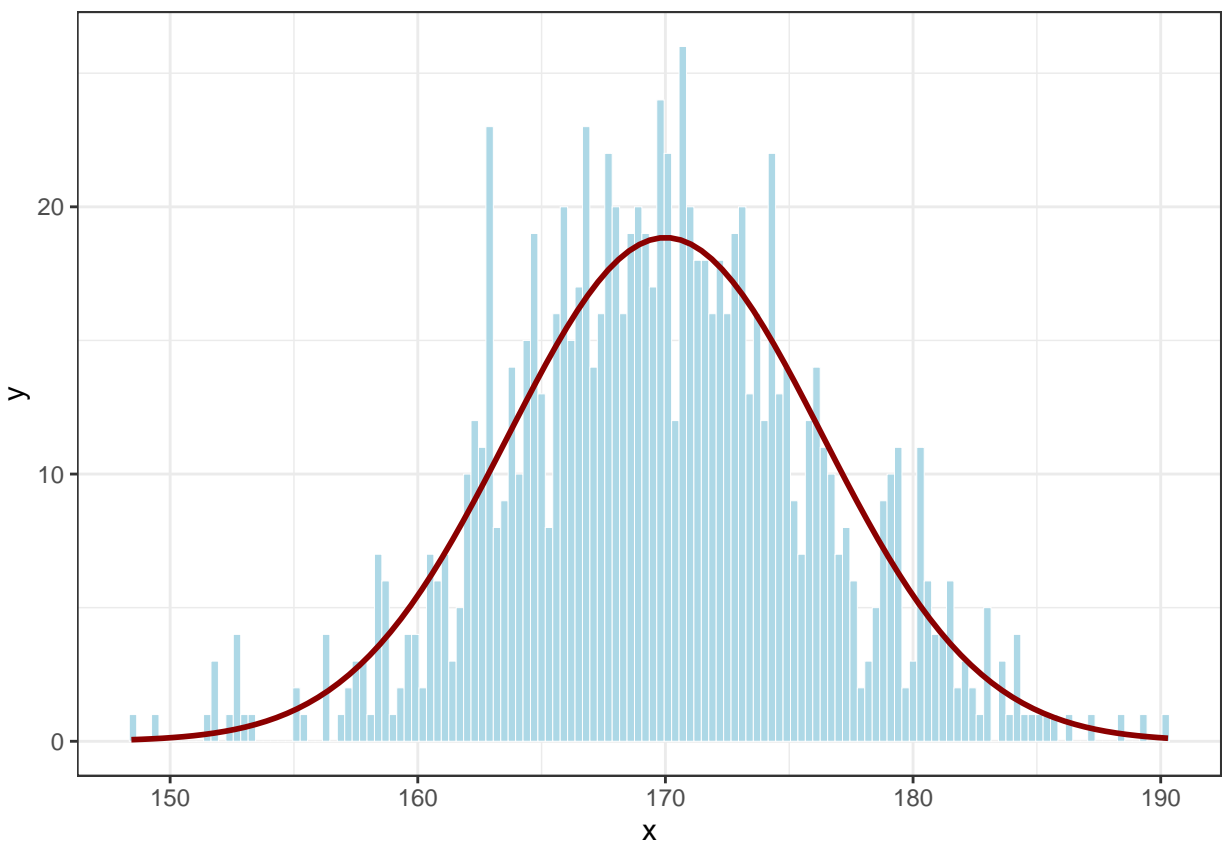
## Normal distribution

Normal distribution, also known as the Gaussian distribution, is a probability distribution that is symmetric about the mean, showing that data near the mean are more frequent in occurrence than data far from the mean.

```

library(tidyverse)
n = 1000
mean = 170 # cm
sd = 6.35 # cm
binwidth= 0.3
set.seed(1234)
df <- data.frame(x=rnorm(n, mean, sd))
ggplot(df, aes(x = x, mean = mean, sd = sd, binwidth = binwidth, n = n))+
  theme_bw()+
  geom_histogram(binwidth = binwidth,
    colour = "white", fill = "lightblue", size = 0.1)+
  stat_function(fun = function(x) dnorm(x, mean = mean, sd = sd) * n * binwidth,
    color = "darkred", size = 1)

```



case when new column

```

library(dplyr)
library(stringr)
feedback <- c('good_book', 'good_read', 'good_story', 'good for knowledge')
book <- c('ramayana', 'bible', 'encyclopedia', 'Mbharatha')

df <- data.frame(feedback, book)

```

```
df %>%  
  mutate(response = case_when(str_starts(feedback, 'good') ~ 'good')) %>%  
  select(book, response) %>% as_tibble()
```

```
## # A tibble: 4 x 2  
##   book      response  
##   <chr>      <chr>  
## 1 ramayana    good  
## 2 bible       good  
## 3 encyclopedia good  
## 4 Mbharatha   good
```