

# Review of R and RStudio

EES 4891-06/5891-01

Bayesian Statistical Methods

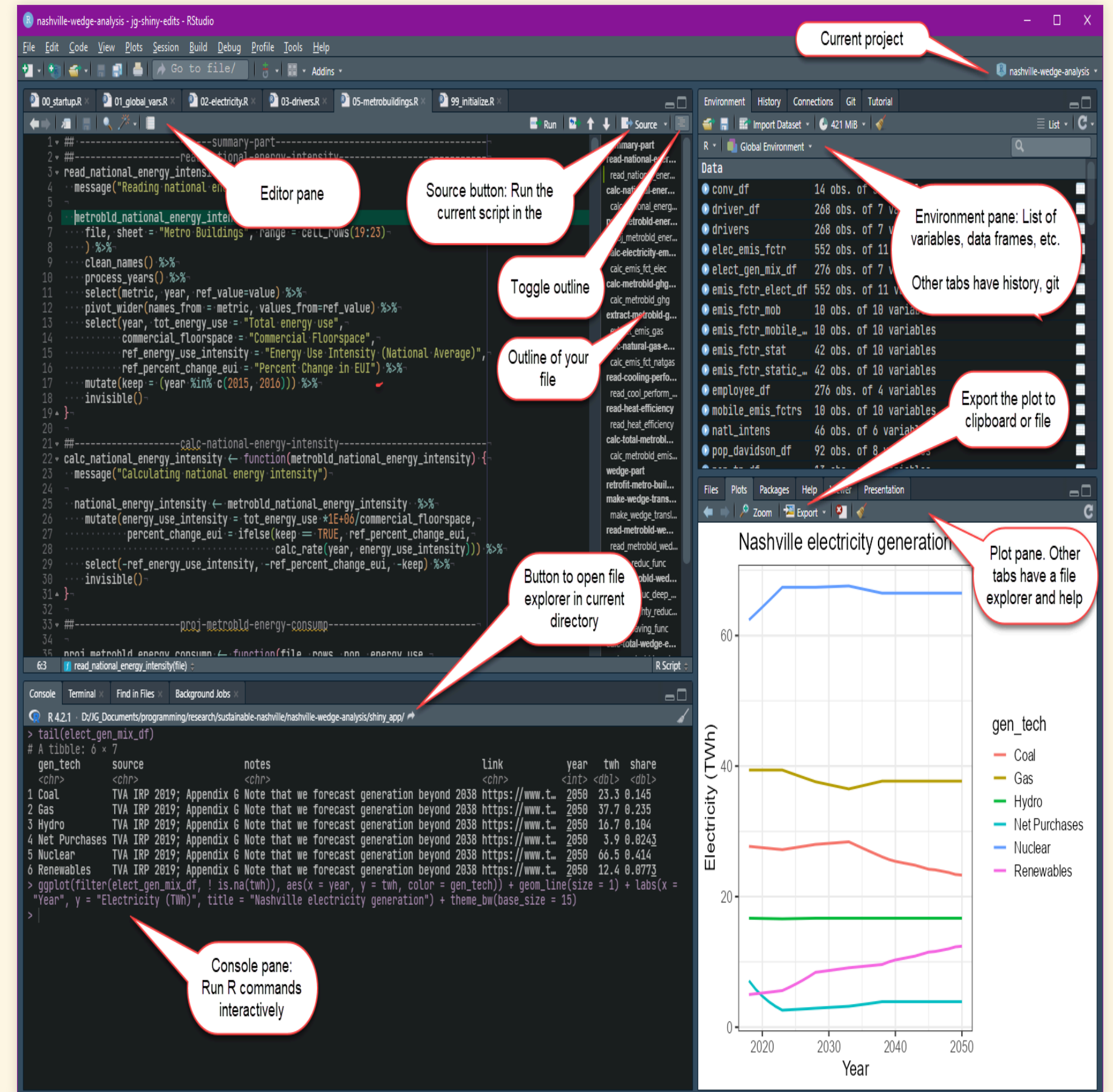
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Class #4: Wednesday, January 14 2026

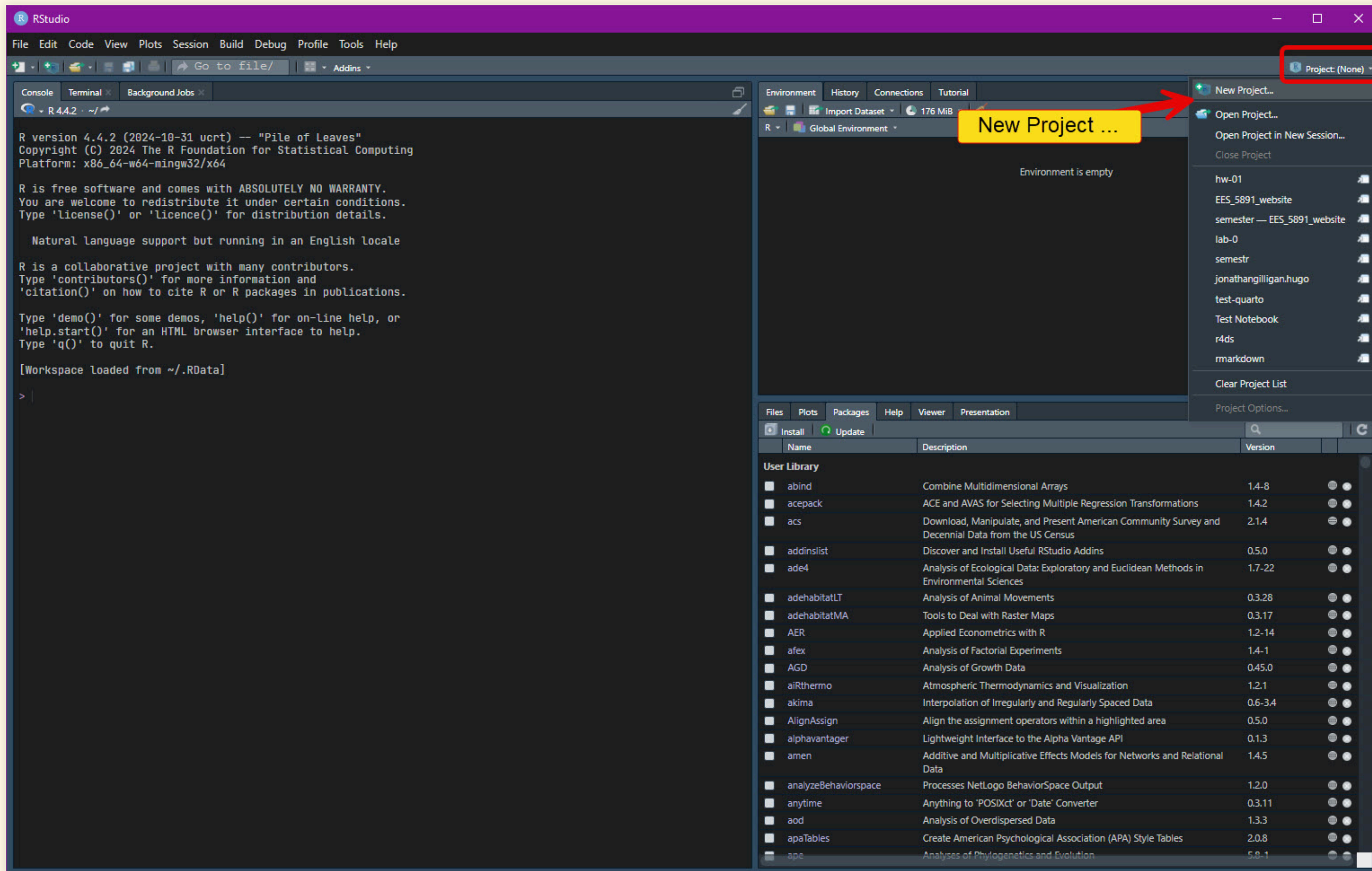
# Overview of Rstudio

# RStudio Window layout

- Project oriented
  - A project is a directory with a group of related files (R, data, etc.)
  - A project can be managed with [git](#) for revision control
  - Create a new project:
    - Create a new directory for a project
    - Create a project in an existing directory
    - Download a project from GitHub or another external source, using [git](#)
- Online help and cheat-sheets
- Window layout
  - Four panes:
    1. Editor pane (edit R scripts, text files, etc.)
    2. Console and Terminal pane (interactively run R commands, etc.)
    3. Files, Plots, Packages, Help, etc.
      - You can export plots to the clipboard or files.
    4. Environment, History, Git, etc.

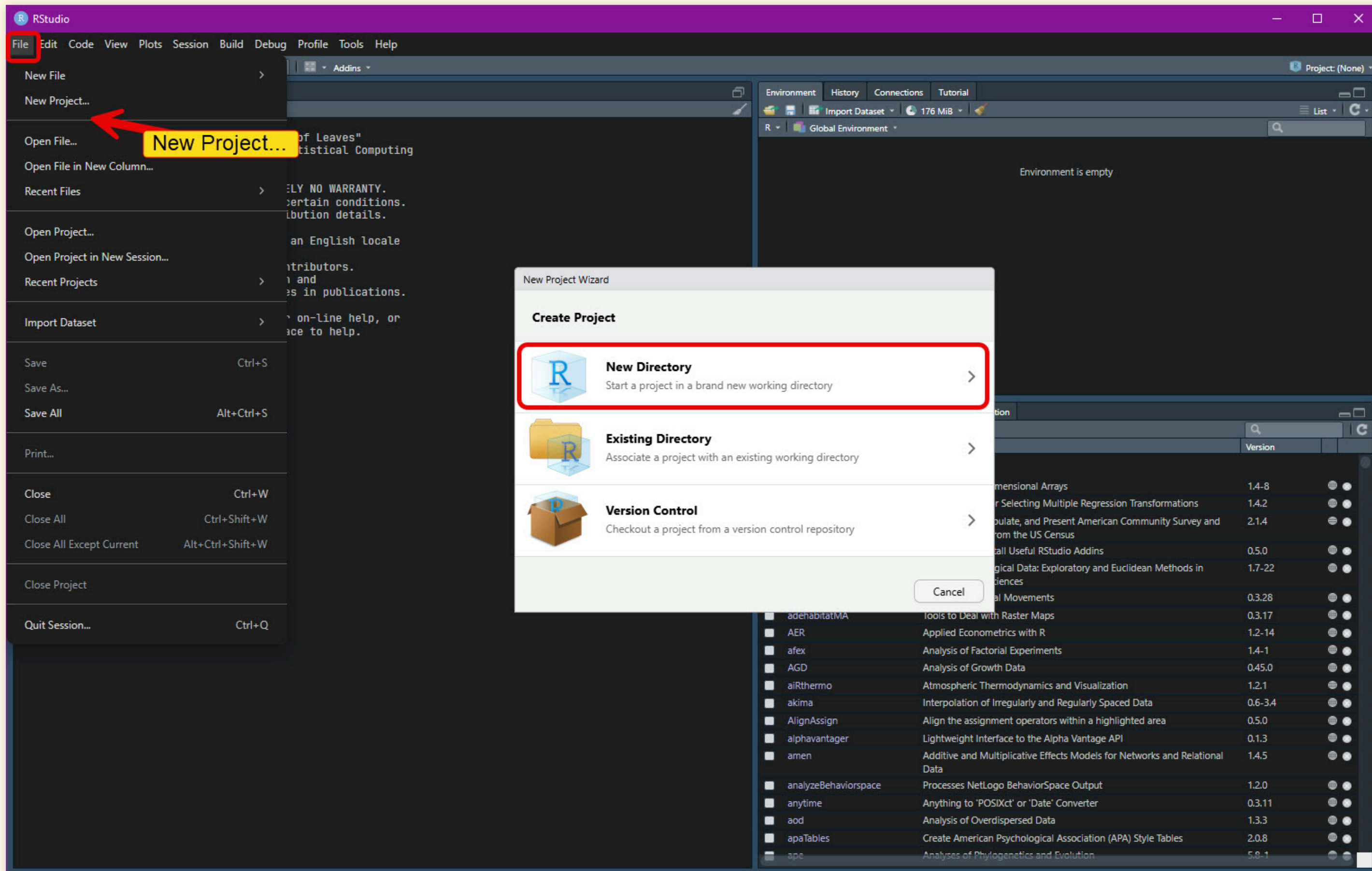


# Creating an RStudio Project





# Creating an RStudio Project



# R Language

# R Language

- R is flexible
- Two approaches
  - Base R
    - Very flexible and powerful
    - Sometimes confusing and verbose
    - Nothing special about data frames (`data.frame`)
  - Tidyverse (`library(tidyverse)`)
    - Unified philosophy of data analysis
      - Canonical reference: [R for Data Science](#)
      - Oriented toward `data.frames` (and `tibbles`)
      - Principles of “Tidy Data”
      - Consistent approach makes it easy to figure out how to do what you want to do.
      - Download extensive cheat-sheets via RStudio help menu.

# Working with Data



- R data types:
  - `integer`, `numeric` (real), `character`, `logical`, `factor`, ...
- R data structures:
  - 1-dim: `vector`, `list`
  - 2-dim: `matrix`, `data.frame`, ...
  - *n*-dim: `array`
  - Data types:
    - `vector`, `array`, `matrix`: elements must be the same type
    - `list`: elements can be different types
  - `data.frame`, `tibble`:
    - Like a spreadsheet or database table.
    - A `list` of `vectors`:
    - Each column is a `vector` representing a variable:
      - height, weight, ...
      - Each column can be a different type.
    - Each row is a set of related measurements for a given person (height, weight, ...).

- Anthropometric data: !Kung San people

```
library(tidyverse)
library(tidybayes)
library(rethinking)
library(tidybayes.rethinking)
```

```
data(Howell11)
d <- tibble(Howell11)
head(d)
```

```
## # A tibble: 6 × 4
##   height weight   age  male
##   <dbl> <dbl> <dbl> <int>
## 1  152.   47.8   63     1
## 2  140.   36.5   63     0
## 3  137.   31.9   65     0
## 4  157.   53.0   41     1
## 5  145.   41.3   51     0
## 6  164.   63.0   35     1
```

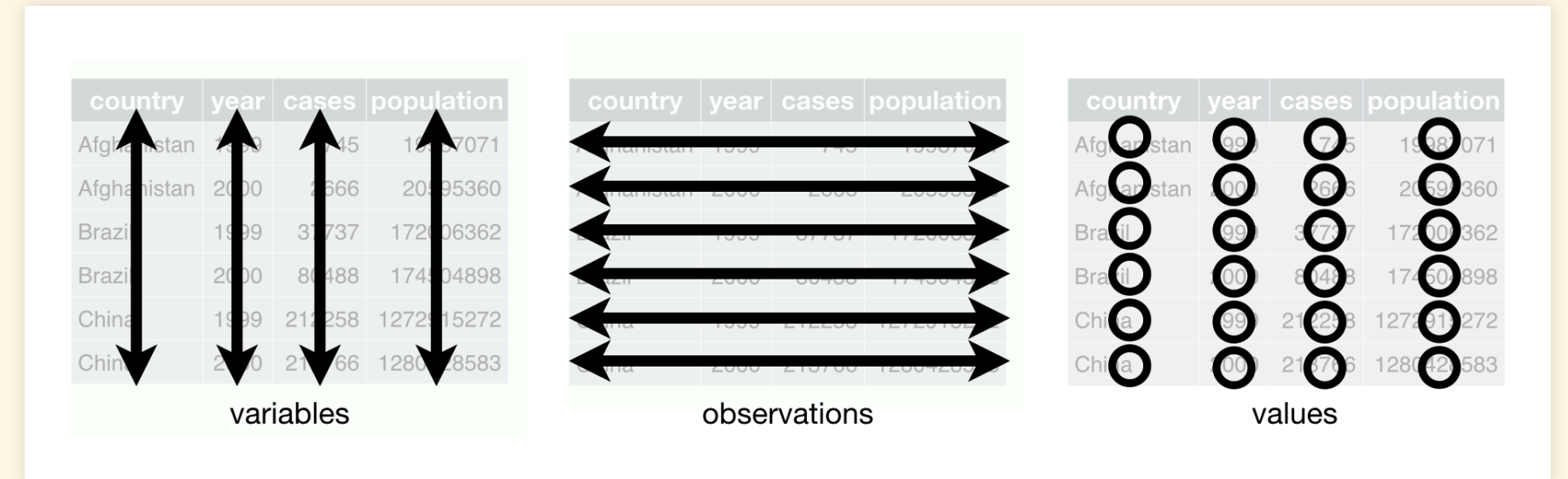
```
precis(d)
```

```
##               mean          sd      5.5%
94.5%
## height 138.2635963 27.6024476 81.108550
165.73500
## weight  35.6106176 14.7191782  9.360721
54.50289
## age      29.3443934 20.7468882  1.000000
66.13500
## male      0.4724265  0.4996986  0.000000
1.00000
##
##           histogram
## height  _____
## weight  _____
## age     _____
## male    _____
```

# Data Structures

# Data Structures

- **Data** is represented as a `data.frame`, or a special kind of data frame called a `tibble` (for “Tidy Table”)
  - A `data.frame` is like a spreadsheet.
  - Each column represents a different variable (year, month, temperature, rainfall, humidity, etc.)
  - Each row represents a different measurement of all the variables.



- Columns can be different data types (integer, numeric, character, logical, factor, ...)
- All rows of a column are the same type

# Data

- Base R

- Separate 1-dimensional lists or vectors of data
- 2D (or higher) arrays: `data.frame`, `matrix`, `array`
- Index rows and columns:
  - Indexing starts at 1
    - Like Matlab, unlike Python, which starts at 0
  - `Howell11[107,]` for all columns of the row
  - `Howell11[, "height"]` for all rows of the "height" column
  - `Howell11[10:15, c(1, 3, 4)]` to get rows 10–15 of columns 1, 3, and 4.
  - `Howell11["age" >= 18, c("height", "weight")]`

- Tidyverse (`library(tidyverse)`)

- `data.frame` or `tibble`

- Select columns:

```
select(Howell11, height, weight, age)
select(Howell11, -male, -age)
```

- Select all columns that start with "foo\_" but don't end with "bar"

```
select(my_data, starts_with("foo_"),
       -ends_with("bar"))
```

- Select rows:

```
filter(Howell11, age >= 18, male)
```

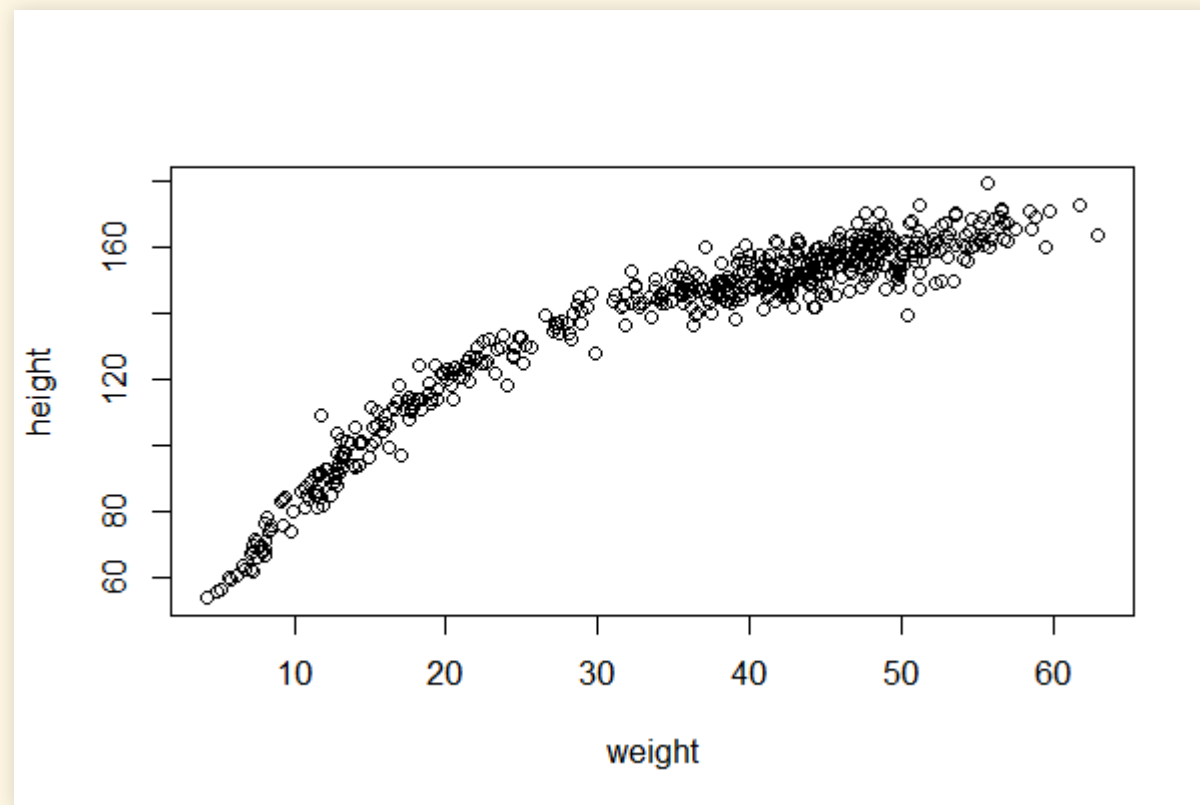
# Graphics

- Base R

- `plot` command

```
plot(x, y) # plot y vs. x with points  
plot(x, y, type = "l") # plot with a line
```

```
plot(height ~ weight, data = Howell1) # with a data.frame
```

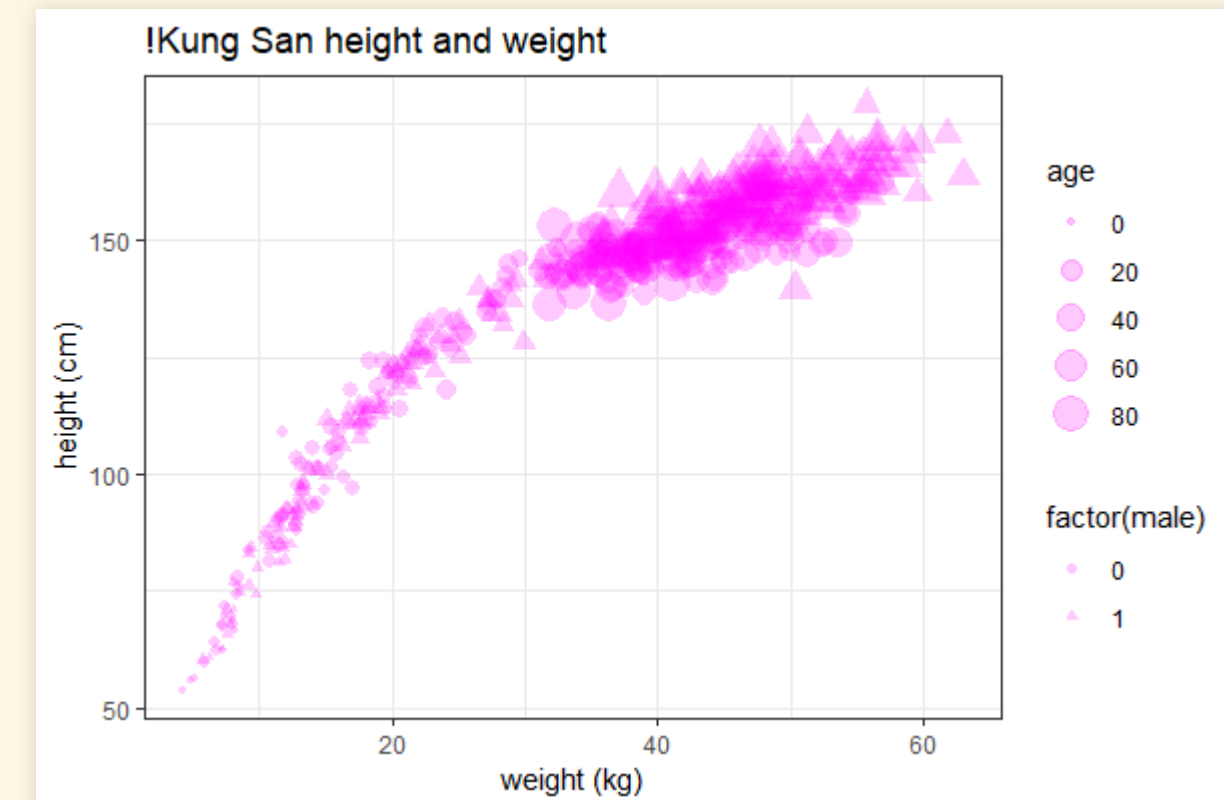


- Tidyverse

- `ggplot` command

```
ggplot(Howell1, aes(x = weight, y = height)) +  
  geom_point() +  
  labs(x = "weight (kg)", y = "height (cm)",  
        title = "!Kung San height and weight")
```

```
ggplot(Howell1, aes(x = weight, y = height)) +  
  geom_point(aes(shape = factor(male), size = age),  
             color = "magenta", alpha=0.2) +  
  labs(x = "weight (kg)", y = "height (cm)",  
        title = "!Kung San height and weight") +  
  theme_bw()
```





# Grammar of Graphics

# Grammar of Graphics

- The heart of data visualization in the Tidyverse is the `ggplot2` package.
  - `gg` from Leland Wilkinson, *The Grammar of Graphics*
- A graph has a grammar, which can be expressed in terms of 7 layers

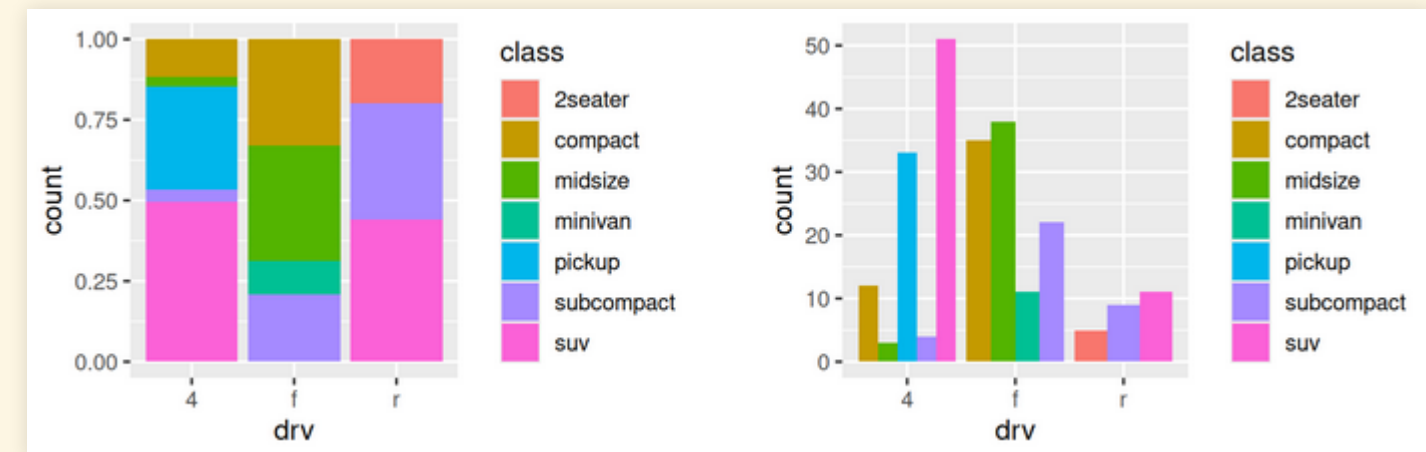
## 1. Data

## 2. Geometries: How to draw the data (Lines, points, etc.)

## 3. Aesthetics: How to mapping different dimensions of the data to visual attributes (x & y coordinates, colors, shape, etc.)

## 4. Statistical transformations: histograms, smoothing, box-plots, etc.

## 5. Position adjustments: e.g., side-by-side versus stacked bar charts



## 6. Coordinate systems: Cartesian vs. polar, pie charts, etc.

## 7. Facets: Breaking a graph into sub-plots.

# Grammar of Graphics in R

- To start a graph, specify the data and aesthetics:

```
ggplot(Howell1, aes(x = weight, y = height))
```

- Next, we add geometries

```
ggplot(Howell1, aes(x = weight, y = height)) +  
geom_point(alpha = 0.2)
```

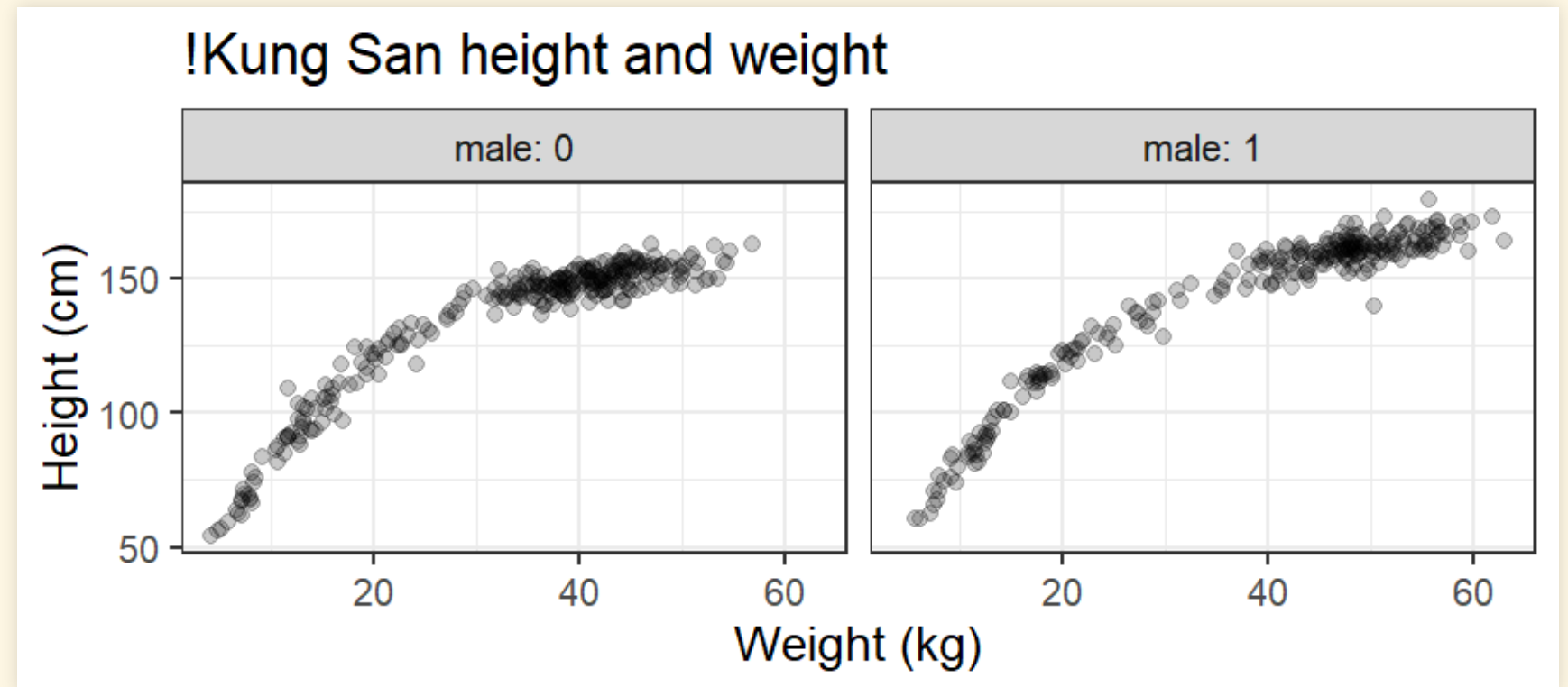
**alpha** makes the points semi-transparent

- Label coordinates

```
ggplot(Howell1, aes(x = weight, y = height)) +  
geom_point(alpha = 0.2) +  
labs(x = "Weight (kg)", y = "Height (cm)",  
      title = "!Kung San height and weight")
```

- Use facets to separate males and females

```
ggplot(Howell1, aes(x = weight, y = height)) +  
geom_point(alpha = 0.2) +  
labs(x = "Weight (kg)", y = "Height (cm)",  
      title = "!Kung San height and weight") +  
facet_wrap(~male, labeller = "label_both")
```



# Jura Data Set

# Jura Data Set



- Survey of soil contamination in the Swiss Jura
  - O. Attela, J.-P. Dubois, & R. Webster. 1994. *Environ. Pollution* **86**, 315.
- The data set contains measurements of cadmium, cobalt, chromium, copper, nickel, lead, and zinc concentrations from 100 soil samples.

```
library(tidyverse)
library(gstat)
data(jura)
jura <- as_tibble(jura.val) |> select(-(Xloc:Yloc))
```

- You may need to run `install.packages("gstat")` in the RStudio console.

```
head(jura)

## # A tibble: 6 × 11
##   long   lat Landuse Rock      Cd    Co    Cr    Cu    Ni    Pb    Zn
##   <dbl> <dbl> <fct>   <fct>   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1  6.85  47.1 Meadow Quaternary 1.57   8.28  37.1 18.6  18.6  38.2  65.2
## 2  6.87  47.2 Meadow Argovian   2.04  10.8  40.8 11.5  21.5  33.4 113.
## 3  6.87  47.2 Pasture Argovian   1.20  12    53.2 13.0  23.9  26.6  91.6
## 4  6.86  47.1 Pasture Quaternary 0.49  10.9  23.4  5.64 14.6  25.9  41.2
## 5  6.84  47.1 Meadow Sequanian 0.692  8.12  27.2 10.3  14.6  31.2  50.4
## 6  6.87  47.1 Forest Kimmeridgian 1.75   9.12  35.5  8.36  26.4  37.7  63.2
```

```
glimpse(jura)

## Rows: 100
## Columns: 11
## $ long      <dbl> 6.854080, 6.865951, 6.871425, 6.857691, 6.837664, 6.871310,
## $ lat       <dbl> 47.14342, 47.1514, 47.15390, 47.13966, 47.13603, 47.13773,
## $ Landuse    <fct> Meadow, Meadow, Pasture, Pasture, Meadow, Forest, Forest,
## $ Rock       <fct> Quaternary, Argovian, Argovian, Quaternary, Sequanian,
## $ Cd         <dbl> 1.570, 2.045, 1.203, 0.490, 0.692, 1.750, 0.415, 0.685,
## $ Co         <dbl> 8.280, 10.800, 12.000, 10.920, 8.120, 9.120, 9.120, 11.720,
## $ Cr         <dbl> 37.12, 40.80, 53.20, 23.40, 27.16, 35.48, 30.32, 31.92,
## $ Cu         <dbl> 18.600, 11.480, 13.040, 5.640, 10.320, 8.360, 4.440, 10.920,
## $ Ni         <dbl> 18.60, 21.52, 23.92, 14.60, 14.64, 26.40, 24.24, 13.12,
## $ Pb         <dbl> 38.20, 33.36, 26.56, 25.88, 31.16, 37.72, 41.00, 30.84,
## $ Zn         <dbl> 65.20, 112.80, 91.60, 41.20, 50.40, 63.20, 53.16, 49.28,
```

# Transforming Rows

- Selecting:

```
filter(jura, Landuse == "Meadow") |> head()
```

```
## # A tibble: 6 × 11
##   long   lat Landuse Rock      Cd    Co    Cr    Cu    Ni    Pb
##   <dbl> <dbl> <fct>   <fct>    <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1  6.85  47.1 Meadow Quaternary 1.57   8.28  37.1  18.6  18.6  38.2
## 2  6.87  47.2 Meadow Argovian   2.04  10.8  40.8  11.5  21.5  33.4
## 3  6.84  47.1 Meadow Sequanian 0.692  8.12  27.2  10.3  14.6  31.2
## 4  6.85  47.1 Meadow Kimmeridgian 0.92  10.6  49.0  30.3  31.5  68.1
## 5  6.85  47.1 Meadow Argovian   0.495  8.52  31.4  17.1  16.1  46.8
## 6  6.84  47.1 Meadow Sequanian 1.19   9.68  37.4  31.4  22.4  72.4
```

- Sorting:

```
filter(jura, Landuse == "Meadow") |>
  arrange(Rock, long, lat) |>
  head()
```

```
## # A tibble: 6 × 11
##   long   lat Landuse Rock      Cd    Co    Cr    Cu    Ni    Pb    Zn
##   <dbl> <dbl> <fct>   <fct>    <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1  6.83  47.1 Meadow Argovian 0.475  3.96  22.2  22.7  7.92  55.2  46.8
## 2  6.84  47.1 Meadow Argovian 3.78   9.68  42.8  32.8  23.5  94.4 175.
## 3  6.84  47.1 Meadow Argovian 0.585  5.8   39.9  15.2  13.2  56.4  51.2
## 4  6.84  47.1 Meadow Argovian 0.57   4.08  24.9  21.4  9.68  67.2  56.8
```

# Transforming Rows

- Selecting on multiple criteria:

```
filter(jura, Landuse == "Meadow", Rock == "Quaternary", Cd > 1.3) |>
  head()
```

```
## # A tibble: 5 × 11
##   long   lat Landuse Rock      Cd    Co    Cr    Cu    Ni    Pb
##   <dbl> <dbl> <fct>   <fct>   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1  6.85  47.1 Meadow Quaternary 1.57  8.28  37.1  18.6  18.6  38.2
## 2  6.86  47.2 Meadow Quaternary 1.58  5.8   40.4  56.4  22.5  93.6 109.
## 3  6.88  47.1 Meadow Quaternary 1.42 11.1   27.5  18.8  20.6  36.5
## 4  6.86  47.1 Meadow Quaternary 2.08 13.2   45.9  39    26.4  52.4 104
## 5  6.88  47.1 Meadow Quaternary 2.61 20.6   37.2  24    29.4  47.2
## 86.4
```

# Transforming Columns





## 1 65.2	6.85	47.1	Meadow	Quaternary	1	0.586	1.57	8.28	37.1	18.6	18.6	38.2
## 2 113.	6.87	47.2	Meadow	Argovian	0.533	0.296	2.04	10.8	40.8	11.5	21.5	33.4
## 3 91.6	6.87	47.2	Pasture	Argovian	0.545	0.290	1.20	12	53.2	13.0	23.9	26.6
## 4 41.2	6.86	47.1	Pasture	Quaternary	0.386	0.628	0.49	10.9	23.4	5.64	14.6	25.9
## 5 50.4	6.84	47.1	Meadow	Sequanian	0.705	0.618	0.692	8.12	27.2	10.3	14.6	31.2
## 6 63.2	6.87	47.1	Forest	Kimmeridgian	0.317	0.597	1.75	9.12	35.5	8.36	26.4	37.7

# Pipes

# Pipes

- The pipe `%>%` or `|>` sends the output from one function into another.
- This code is hard to read:

```
relocate( rename( mutate( jura,  
                           CuNi = Cu / Ni,  
                           PbZn = Pb/Zn),  
             longitude = long,  
             latitude = lat),  
          CuNi:PbZn, .before = Cd)
```

- This code is much easier to read

```
jura |>  
  mutate(CuNi = Cu / Ni, PbZn = Pb/Zn) |>  
  rename(longitude = long, latitude = lat) |>  
  relocate(CuNi:PbZn, .before = Cd)
```

- We could also write

```
tmp <- mutate(jura, CuNi = Cu / Ni, PbZn = Pb/Zn)  
tmp <- rename(tmp, longitude = long,  
              latitude = lat)  
relocate(tmp, CuNi:PbZn, .before = Cd)
```

# Summarizing and Grouping



# Summarizing and Grouping

- Average lead content:

```
jura |> summarize(mean_Pb = mean(Pb),  
                  sd_Pb = sd(Pb),  
                  count = n())
```

```
## # A tibble: 1 × 3  
##   mean_Pb sd_Pb count  
##   <dbl> <dbl> <int>  
## 1   56.5  40.5   100
```

- Grouped Summaries:

```
jura |> group_by(Landuse, Rock) |>  
  summarize(mean_Pb = mean(Pb),  
            sd_Pb = sd(Pb), count = n()) |>  
  ungroup()
```

```
## # A tibble: 15 × 5  
##   Landuse Rock      mean_Pb sd_Pb count  
##   <fct>   <fct>      <dbl> <dbl> <int>  
## 1 Forest Argovian      33.9  3.21     4  
## 2 Forest Kimmeridgian  53.6 15.3    10  
## 3 Forest Sequanian    57.9 13.4     2  
## 4 Forest Portlandian  42.0  9.56     2  
## 5 Pasture Argovian    28.5 11.4     4  
## 6 Pasture Kimmeridgian 60.9 69.4    14  
## 7 Pasture Sequanian   42.1 11.5     6  
## 8 Pasture Quaternary  37.1 15.9     2  
## 9 Meadow Argovian    53.1 25.3    14  
## 10 Meadow Kimmeridgian 50.1 11.9    13  
## 11 Meadow Sequanian   73.1 54.1    18  
## 12 Meadow Portlandian 109.  NA      1  
## 13 Meadow Quaternary  60.8 41.5     7  
## 14 Tillage Argovian   153.  NA      1  
## 15 Tillage Kimmeridgian 42.3  5.77     2
```

# More Grouping

- Counting combinations:

```
jura |> count(Landuse, Rock)
```

```
## # A tibble: 15 × 3
##   Landuse Rock      n
##   <fct>   <fct>   <int>
## 1 Forest  Argovian     4
## 2 Forest  Kimmeridgian 10
## 3 Forest  Sequanian    2
## 4 Forest  Portlandian   2
## 5 Pasture Argovian     4
## 6 Pasture Kimmeridgian 14
## 7 Pasture Sequanian    6
## 8 Pasture Quaternary    2
## 9 Meadow  Argovian    14
##10 Meadow  Kimmeridgian 13
##11 Meadow  Sequanian   18
##12 Meadow  Portlandian   1
##13 Meadow  Quaternary    7
##14 Tillage Argovian     1
##15 Tillage Kimmeridgian 2
```

- Alternate Grouping:

```
jura |> summarize(mean_Pb = mean(Pb),
                  sd_Pb = sd(Pb),
                  count = n(),
                  .by = c("Landuse", "Rock"))
```

```
## # A tibble: 15 × 5
##   Landuse Rock      mean_Pb sd_Pb count
##   <fct>   <fct>      <dbl> <dbl> <int>
## 1 Meadow  Quaternary    60.8  41.5     7
## 2 Meadow  Argovian     53.1  25.3    14
## 3 Pasture Argovian     28.5  11.4     4
## 4 Pasture Quaternary    37.1  15.9     2
## 5 Meadow  Sequanian    73.1  54.1    18
## 6 Forest  Kimmeridgian   53.6  15.3    10
## 7 Pasture Sequanian    42.1  11.5     6
## 8 Meadow  Kimmeridgian   50.1  11.9    13
## 9 Pasture Kimmeridgian   60.9  69.4    14
##10 Forest  Portlandian    42.0   9.56     2
##11 Forest  Argovian     33.9   3.21     4
##12 Meadow  Portlandian   109.   NA        1
##13 Tillage Kimmeridgian   42.3   5.77     2
##14 Forest  Sequanian    57.9  13.4     2
##15 Tillage Argovian    153.   NA        1
```

# Slicing

- Selecting:

```
jura |> group_by(Landuse, Rock) |>
  slice_max(Pb, n = 1)
```

##	#	A tibble: 15 × 11									
##	#	Groups: Landuse, Rock [15]									
##		long	lat	Landuse	Rock	Cd	Co	Cr	Cu	Ni	
Pb		Zn									
##		<dbl>	<dbl>	<fct>	<fct>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	
<dbl>	<dbl>										
##	1	6.86	47.2	Forest	Argovian	1.32	3.74	27.6	5.4	14.4	
37.5		46.4									
##	2	6.86	47.1	Forest	Kimmeridgian	1.25	8.08	39.6	13.1	18.6	88
86.8											
##	3	6.85	47.1	Forest	Sequanian	1.01	9.96	28.7	5.96	17.4	
67.4		52.5									
##	4	6.87	47.1	Forest	Portlandian	1.22	5.24	27.0	5.52	21.0	
48.8		46.4									
##	5	6.85	47.1	Pasture	Argovian	0.375	12.0	34.1	19.4	16.4	
45.2		70									
##	6	6.86	47.1	Pasture	Kimmeridgian	1.76	10.3	40.5	127	30.8	300
192											
##	7	6.83	47.1	Pasture	Sequanian	2.54	12.6	70	8.72	26.2	
55.6		71.6									
##	8	6.88	47.1	Pasture	Quaternary	1.31	12.7	34.8	17.7	19.6	
48.4		80.4									
##	9	6.85	47.1	Meadow	Argovian	0.394	4.44	21.6	39.6	8.92	106.
72.4											
##	10	6.85	47.1	Meadow	Kimmeridgian	0.825	15.3	36.5	31.2	25.4	
70.4		75.4									
##	11	6.86	47.1	Meadow	Sequanian	1.78	11.4	41	155.	24.5	240.
260.											
##	12	6.88	47.1	Meadow	Portlandian	1.62	12.0	34.6	91.2	30.2	109.
157.											
##	13	6.85	47.1	Meadow	Quaternary	0.75	15.6	29.8	73.1	20.2	139.
95.7											
##	14	6.85	47.1	Tillage	Argovian	1.31	8.44	41.6	118.	20.4	153.
145.											
##	15	6.87	47.1	Tillage	Kimmeridgian	1.93	13.8	45	19.3	35.7	
46.4		90									

- Random Sampling

```
jura |> slice_sample(n = 5)
```

```
## # A tibble: 5 × 11
##   long   lat Landuse Rock      Cd    Co    Cr    Cu    Ni    Pb
Zn
##   <dbl> <dbl> <fct>   <fct>      <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
<dbl>
## 1  6.83  47.1 Meadow  Argovian    0.475  3.96  22.2  22.7   7.92  55.2
46.8
## 2  6.84  47.1 Meadow  Argovian    3.78  9.68  42.8  32.8  23.5  94.4
175.
## 3  6.86  47.1 Meadow  Argovian    0.745  3.95  26.6   8.52  15.4  30.3
52.4
## 4  6.86  47.1 Meadow  Kimmeridgian 1.68  14.4  46.3  22.9  43.7  61.4
111.
## 5  6.85  47.1 Meadow  Quaternary   1.57   8.28  37.1  18.6  18.6  38.2
65.2
```

# Sampling from models

- Base R

```
library(rethinking)

data(Howell1)
adults <- filter(Howell1, age >= 18)

mdl <- quap(alist(
  height ~ dnorm(mu, sigma),
  mu <- a + b * weight,
  a ~ dnorm(178, 20),
  b ~ dlnorm(0, 1),
  sigma ~ dunif(0, 50)
), data = adults)

w_lst <- data.frame(weight = seq(30, 70, by = 5))
```

- Sample from the posterior predictive distribution for data (e.g., [height](#)):

```
extract.samples(mdl, 1000)
```

- Sample from the posterior of the model link ([mu](#)):

```
link(mdl, w_lst)
```

- Tidyverse ([library\(tidyverse\)](#))

```
library(tidybayes)
library(tidybayes.rethinking)
```

- Sample from the posterior predictive distribution for data (e.g., [height](#)):

```
predicted_draws(mdl, w_lst, ndraws = 1000,
                 value = "height")
add_predicted_draws(w_lst, mdl, ndraws = 1000,
                    value = "height")
```

- Sample from the posterior of the model link ([mu](#)):

```
linpred_draws(mdl, w_lst, ndraws = 1000,
               value = "height")
add_linpred_draws(w_lst, mdl, ndraws = 1000,
                  value = "height")
```

- [add\\_](#) versions are the same, but reverse the order of the first two arguments (model object and new data).

