

Lists in R

Data structure: a structure, usually with a specific class, that holds data.
eg vector, matrix, array, list, tibble

We are already familiar somewhat with vectors

Vectors are a type of data structure.

`fib ← c(1, 2, 3, 5, 8, 13)`

^
all values in the list
have the same class
`class(fib)` is ?

`she_loves_me_she_loves_me_not ←`

`c(TRUE, FALSE, TRUE, FALSE, TRUE)`

`word_on_street ← c("I", "am", "the", "king")`

Vectors are very important in R.

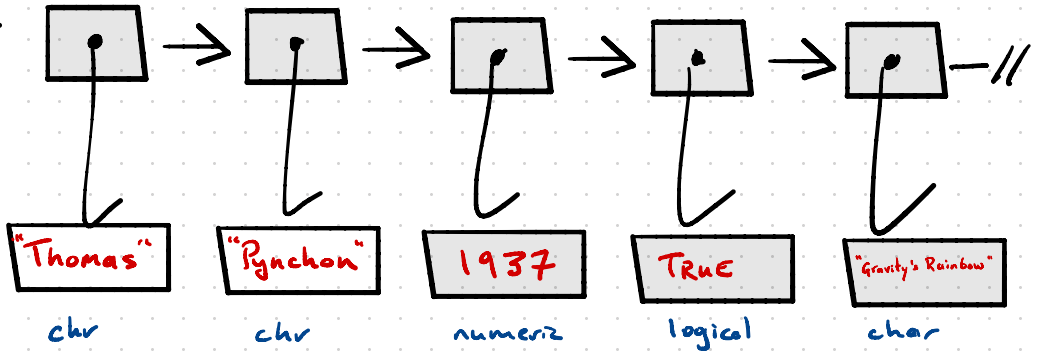
Think of one column (variable) in a tibble.

Lists are more general, more flexible.

lists are another type of data structure.

$L \leftarrow \text{list}(\text{"Thomas"}, \text{"Pynchon"}, 1937, \text{TRUE}, \text{"Gravity's Rainbow"})$

L^2



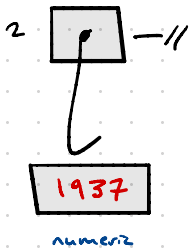
$\text{class}(L)$; $\text{length}(L)$; $\text{names}(L)$

list

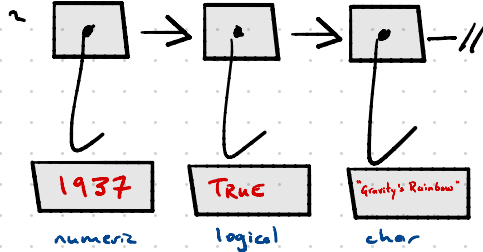
5

NULL

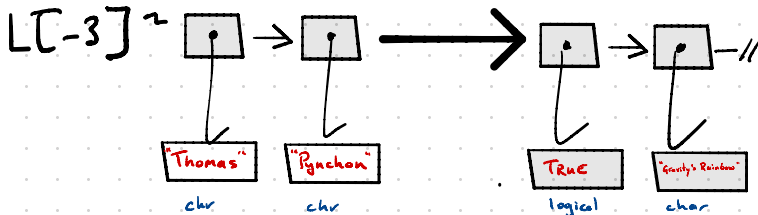
$L[3]$



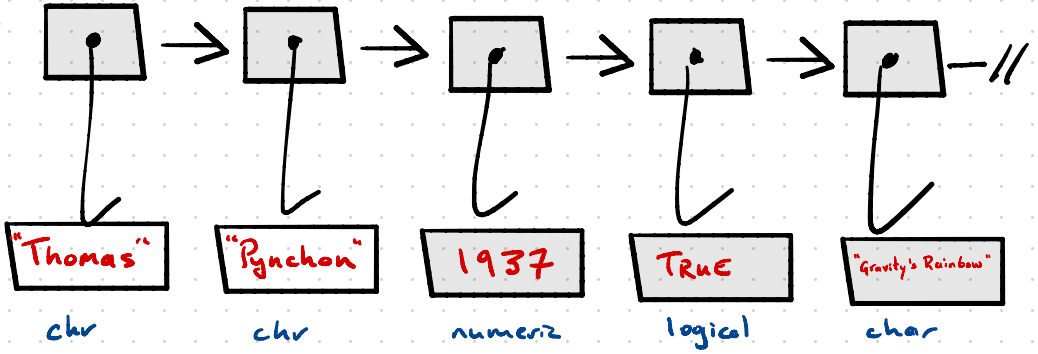
$L[3:5]$



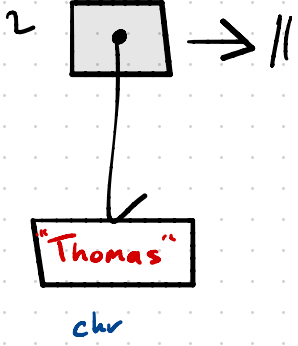
$\text{class}(L[-3]) = ???$



L2

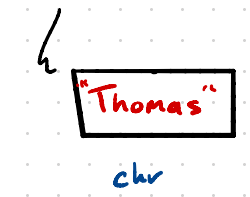


L[1]



THIS IS STILL A LIST (WITH JUST ONE ITEM)

L[[1]]



THIS IS THE VALUE IN THE LIST ^Thomas of class chr.

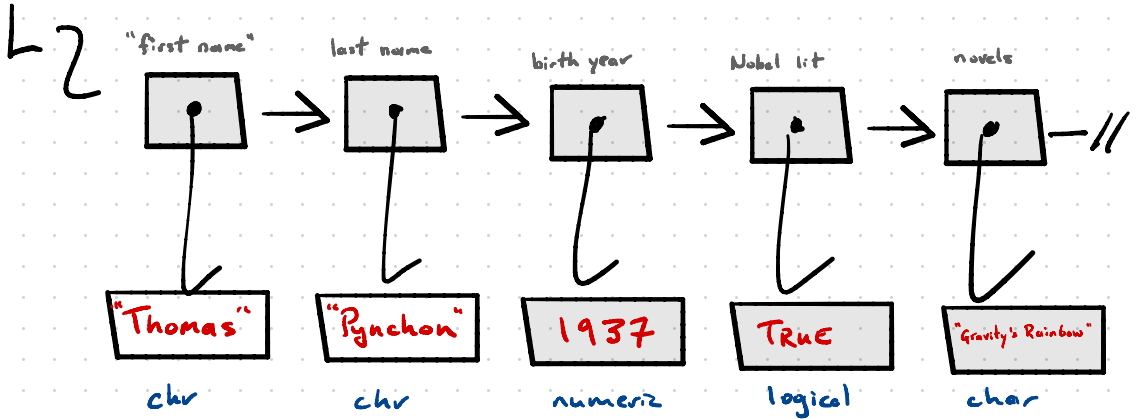
L[[1:3]]



- L[[1]]
- L[[2]]
- L[[3]]

Naming Elements of Lists

names(L) ← c("first name", "last name", "birth year", "Nobel Lit", "novels")



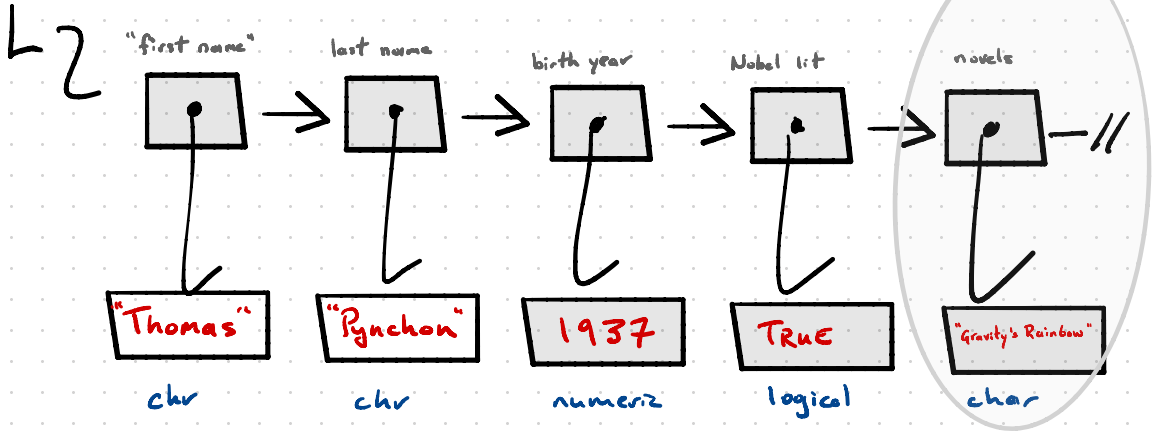
L[3] "birth year"

same as L[[3]]

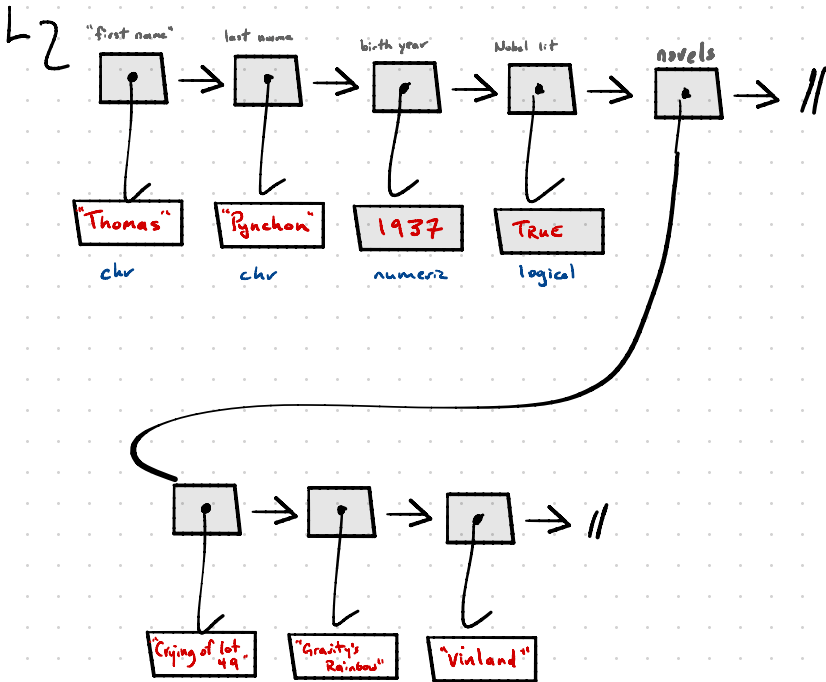
L[5] "novels"

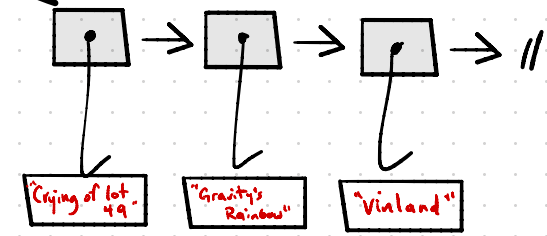
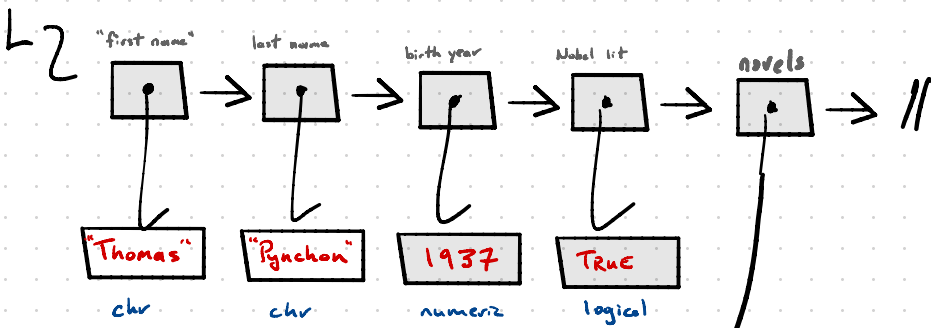
same as L[[5]]

Pynchon has written many novels?



lists can even contain lists.

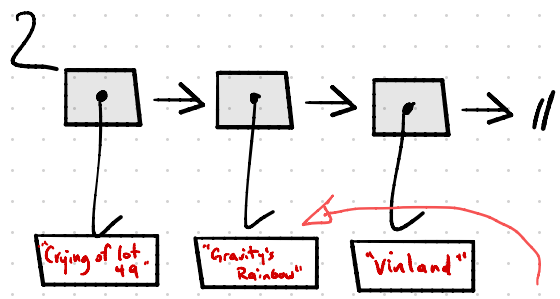




L[[5]]

L \$ novels

L \$ "novels" ?

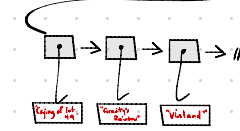
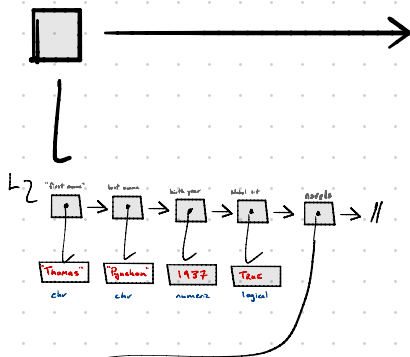


L \$ novels [2:3] ?

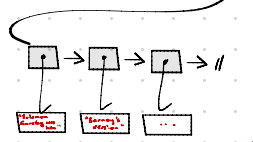
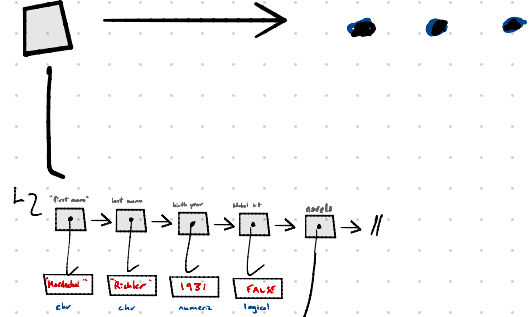
L \$ novels [[2]] ?

Literature

2



Thomas
Pynchon



Mordechai
Richler.

DAVID SANKOFF

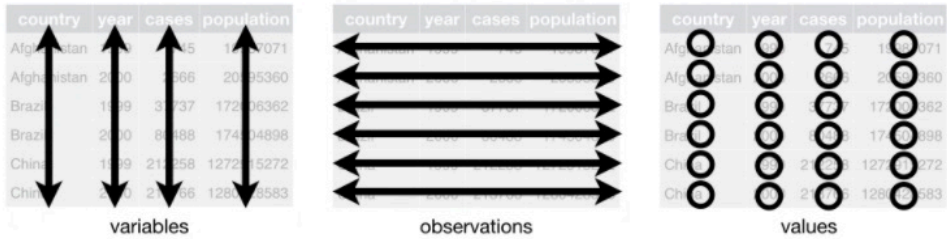
U de M
U de Ottawa.

Tidy Data

There are three interrelated rules which make a dataset tidy:

1. Each variable must have its own column.
2. Each observation must have its own row.
3. Each value must have its own cell.

Figure 12.1 shows the rules visually.



This means for most real analyses, you'll need to do some tidying. The first step is always to figure out what the variables and observations are. Sometimes this is easy; other times you'll need to consult with the people who originally generated the data. The second step is to resolve one of two common problems:

1. One variable might be spread across multiple columns.
2. One observation might be scattered across multiple rows.

Action Item:

Find examples in the Tara Oceans or TCGA data.

ESR1 is one variable but it has 2 columns.

LONGER

①

participant	at diagnosis ESR1	after treatment ESR1
AINF	628	527
⋮	⋮	⋮
⋮	⋮	⋮

T %>%

`pivot_longer(c("at diagnosis ESR1", "after treatment ESR1"),`

`names_to = "measurement point",`

`values_to = "expression"`

fibble ~

participant	at diagnosis ESR1	after treatment ESR1
AINF	628	527
⋮	⋮	⋮



part.	measurement pt.	expression
AINF	at diagnosis ESR	628
AINF	after treatment	527

②

WIDER

One obs. across many rows.

T₂

participant	gene	count
AINF	ESRI	628
AINE	ANLN	527

↗
2 rows

AINF IS ONE OBSERVATION (FOR WHICH WE MEASURE MANY VARIABLES).

T % > %

pivot_wider(names_from = gene, values_from = count)



parti	ESRI	ANLN
AINF	628	527

separate

- when a single variable has >1 pieces of information (values).

Station identifier [TARA_station#]	Date/Time [yyyy-mm-ddThh:mm]	Latitude [degrees North]	Lon [deg East]
TARA_004	2009-09-15 18:00:00	36.5533	
TARA_004	2009-09-15 11:30:00	36.5533	
TARA_007	2009-09-23 16:08:00	37.0541	
TARA_007	2009-09-23 12:50:00	37.0510	
TARA_009	2009-09-28 16:59:00	39.0609	
TARA_009	2009-09-28 12:18:00	39.1633	

Showing 1 to 7 of 243 entries, 17 total columns

Console Terminal Jobs

```
# www.marinerregions.com]` <chr>
>
> View(T2)
> T2$`Date/Time [yyyy-mm-ddThh:mm]`
[1] "2009-09-15 18:00:00 UTC" "2009-09-15 11:30:00 UTC"
[3] "2009-09-23 16:08:00 UTC" "2009-09-23 12:50:00 UTC"
[5] "2009-09-28 16:59:00 UTC" "2009-09-28 12:18:00 UTC"
[7] "2009-11-02 14:07:00 UTC" "2009-11-02 14:07:00 UTC"
[9] "2009-11-02 08:13:00 UTC" "2009-11-02 08:13:00 UTC"
[11] "2009-11-15 00:16:00 UTC" "2009-11-15 12:50:00 UTC"
```

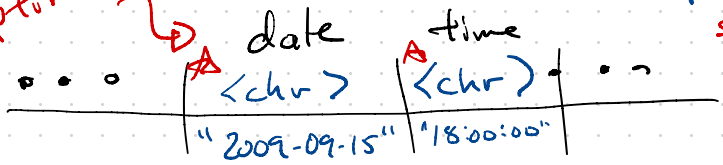


T1 %>%

separate('Date/Time...', into = c("date", "time"),

sep = " ")

not optimal



Tidy data should all be in the same tibble.

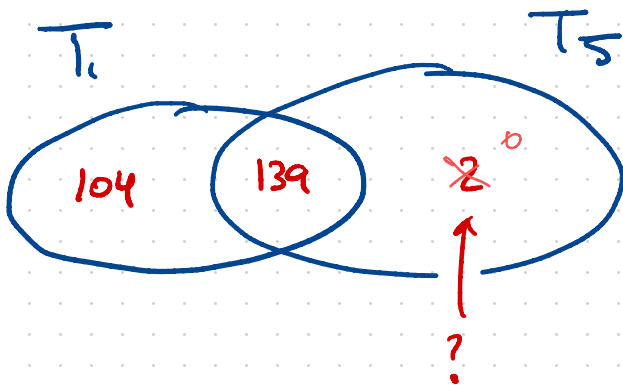
```
> T1
# A tibble: 243 x 18
  sample_label ac_sample ac_run eng pangea_id pangea_data station date      time latitude longitude depth feature
<chr>         <chr>      <chr> <chr> <chr> <chr> <chr> <date> <time> <dbl> <dbl> <dbl> <chr>
1 TARA_004_DC... ERS4487936 ERR59... http... TARA_X00... http://ww... TARA_0... 2009-09-15 18:00 36.6 -6.57 40 (DCM) ...
2 TARA_004_SR... ERS487899 ERR59... http... TARA_Y20... http://ww... TARA_0... 2009-09-15 11:30 36.6 -6.57 5 (SRF) ...
3 TARA_007_DC... ERS477953 ERR31... http... TARA_A20... http://ww... TARA_0... 2009-09-23 16:08 37.1 1.95 42 (DCM) ...
4 TARA_007_SR... ERS477931 ERR31... http... TARA_A20... http://ww... TARA_0... 2009-09-23 12:50 37.1 1.94 5 (SRF) ...
5 TARA_009_DC... ERS488147 ERR59... http... TARA_X00... http://ww... TARA_0... 2009-09-28 16:59 39.1 5.94 55 (DCM) ...
6 TARA_009_SR... ERS488119 ERR59... http... TARA_X00... http://ww... TARA_0... 2009-09-28 12:18 39.2 5.92 5 (SRF) ...
7 TARA_018_DC... ERS488346 ERR59... http... TARA_S20... http://ww... TARA_0... 2009-11-02 14:07 35.8 14.3 60 (DCM) ...
8 TARA_018_DC... ERS488354 ERR59... http... TARA_A10... http://ww... TARA_0... 2009-11-02 14:07 35.8 14.3 60 (DCM) ...
9 TARA_018_SR... ERS488330 ERR59... http... TARA_A10... http://ww... TARA_0... 2009-11-02 08:13 35.8 14.3 5 (SRF) ...
10 TARA_018_SR... ERS488340 ERR59... http... TARA_A10... http://ww... TARA_0... 2009-11-02 08:13 35.8 14.3 5 (SRF) ...
# ... with 233 more rows, and 5 more variables: size_lower <chr>, size_upper <dbl>, pelagic_biomes <chr>, regions <chr>,
# 'pelagic_biomes MRGID' <chr>
> T5
# A tibble: 10 x 5
  pangea_id `16S_mITAGs` richness chao1 shannon_diversity
<chr> <dbl> <dbl> <dbl> <dbl>
1 TARA_B100000965 103040 3053. 4404. 6.83
2 TARA_B100000959 111240 3172. 4805. 6.68
3 TARA_B100000963 122482 2412. 3504. 6.60
4 TARA_B100000902 112336 2931. 4241. 6.78
5 TARA_B100000953 56449 3108. 4723. 6.68
6 TARA_B100000900 74287 2293. 3198. 6.62
7 TARA_B100000927 77265 2501. 3567. 6.68
8 TARA_B100000929 110154 3372. 4999. 6.83
9 TARA_B100000925 128823 2356. 3372. 6.65
10 TARA_B100001113 128677 2874. 4207. 6.80
# ... with 131 more rows
> T8
# A tibble: 245 x 37
  pangea_id data latitude longitude depth `temp[C]` `salinity[PSU]` `oxygen[umol/kg]` `Mean_Nitrates[...
<chr> <dtm> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
1 TARA_B10... 2011-04-15 13:10:00 -13.0 -96.0 57.6 20.6 35.5 217. 1.50
2 TARA_B10... 2011-04-16 16:09:00 -12.9 -96.1 175. 13.0 34.8 0.708 21.1
3 TARA_B10... 2011-04-15 13:10:00 -13.0 -96.0 5.48 25.3 35.8 200. 4.59
4 TARA_B10... 2011-04-22 19:51:00 -5.27 -85.2 45.7 19.6 34.9 104. 20.3
5 TARA_B10... 2011-04-22 14:16:00 -5.27 -85.2 476. 9.20 34.7 4.43 40.0
6 TARA_B10... 2011-04-21 20:16:00 -5.25 -85.2 5.48 24.9 34.7 206. 11.7
7 TARA_R10... 2011-05-13 00:00:00 2.08 -84.5 29.8 26.5 34.3 203. 3.89
8 TARA_B10... 2011-05-13 00:00:00 2.08 -84.5 29.8 26.5 34.3 203. 3.89
9 TARA_B10... 2011-05-13 19:44:00 2.07 -84.5 376. 11.3 34.8 2.52 33.9
10 TARA_R10... 2011-05-12 13:38:00 1.99 -84.6 5.40 27.6 33.4 199. 1.21
# ... with 235 more rows, and 28 more variables: `NO2 [umol/L]**` <dbl>, `P04 [umol/L]**` <dbl>, `NO2NO3 [umol/L]**` <dbl>,
# `SI [umol/L]**` <dbl>, `AMODIS:PAR8d,Einsteins/m-2/d-1` <dbl>, `Okubo-Weiss` <dbl>, `Lyapunov_exp` <dbl>,
# `grad_SST_adv` <dbl>, `retention` <dbl>, `Mean Depth MLD Sigma [m]**` <dbl>, `Mean Depth Max Fluo [m]**` <dbl>, `Mean Depth
# Max N2 [m]**` <dbl>, `Mean Depth Max O2 [m]**` <dbl>, `Mean Depth Min O2 [m]**` <dbl>, `Mean Depth Nitracline
# [m]**` <dbl>, `miTAG.SILVA.Taxo.Richness` <dbl>, `miTAG.SILVA.Phylo.Diversity` <dbl>, `miTAG.SILVA.Chao` <dbl>,
# `miTAG.SILVA.ace` <dbl>, `miTAG.SILVA.Shannon` <dbl>, `OG.Shannon` <dbl>, `OG.Richness` <dbl>, `OG.Evenness` <dbl>, `FC -
# heterotrophs [cells/mL]` <dbl>, `FC - autotrophs [cells/mL]` <dbl>, `FC - bacteria [cells/mL]` <dbl>, `FC -
# picoeukaryotes [cells/mL]` <dbl>, `minimum generation time [h]` <dbl>
```

Quite straight forward to link this data.

pangea-id is both a foreign & primary key. (Section 13.3 keys)

Start with Tables T_i & T_s .

T_i			T_s		
pangea-id	sample-label	...	pangea-id	16S-nr	...
TARA_X20...	TARA_B1...
⋮	⋮	⋮	⋮	⋮	⋮
⋮	⋮	⋮	⋮	⋮	⋮



In general ...

T_i	key	value	...
	T0	5	...
	T1	7	
	T2	3	

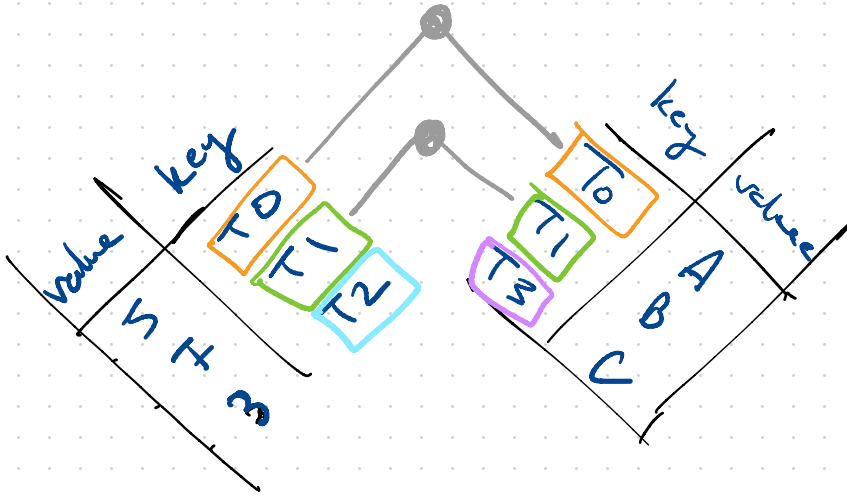
T_s	key	value	...
	T0	A	
	T1	B	..
	T3	C	

T_1

key	value	...
T0	5	..
T1	7	
T2	3	

T_5

key	value
T0	A	
T1	B	..
T3	C	



Inner join:

like an "and" operation

key must be in both tibbles.

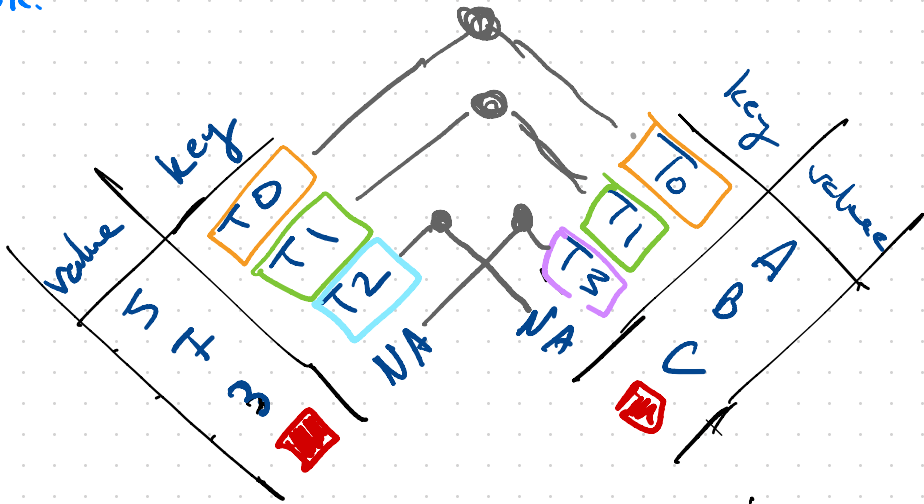
key	value $_{T_1}$	value $_{T_5}$
T0	5	A
T1	7	B

Outer joins: keeps observations (rows) in at least one of the tables.

3 types:

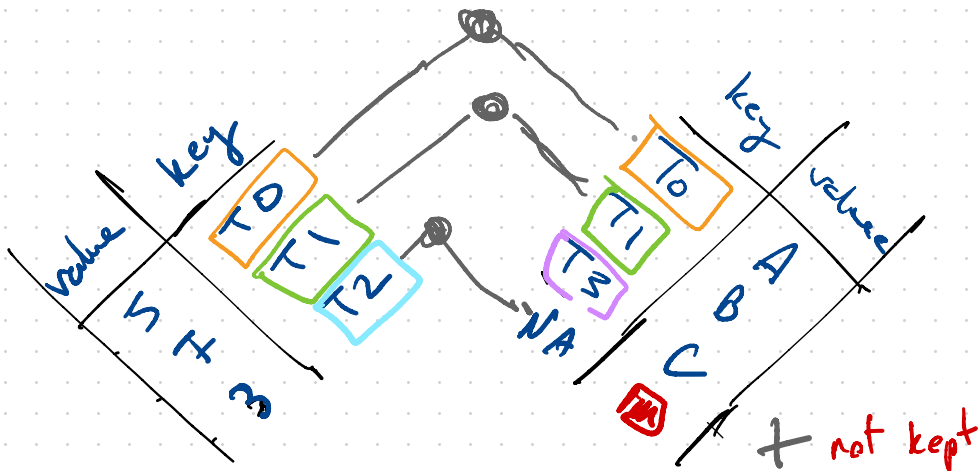
Full join: keeps all observations

↑
sort of like an OR.



key	value T_1	value T_2
T ₀	5	A
T ₁	7	B
T ₂	3	NA
T ₃	NA	C

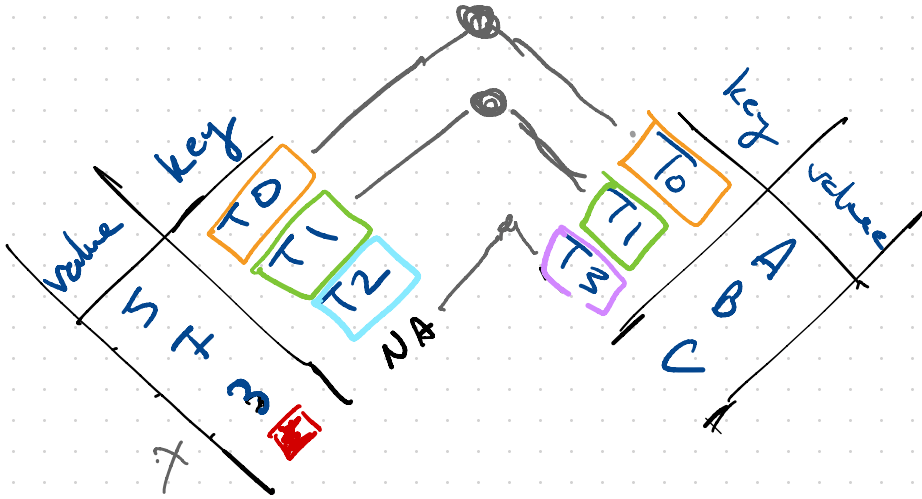
Left join keep everything in first (left) table



key	value _{T1}	value _{T5}
T0	5	A
T1	7	B
T2	3	NA

Right join

keep everything in
table



key	value _{T1}	value _{T3}
T0	5	A
T1	7	B
T3	NA	C

16S rRNA miTAG sequences : 500 Mb.

One sample file. What variable in T does this correspond to?

```
TARA_052_DCM_0.22-1.6.16SrRNA.miTAG.fna
->SOUFRE_0091:3:1305:8579:68103#TGACCA/1
CAGCAGTGGGGAATCTTGACCAATGGGGCCGAAAGCCTGATCCAGCCATTCGCGTGGATGATGAAGGCCCTAGGGTTGTAATAAATCTTTCGGGA
GGGAAGATAATGACGGTACTGCTAAAGAAGCCCGGGCTAACTCCGTGCCAGCAGCCGCGTAATACGGAGGGGGCTAGCGTTG
>SOUFRE_0091:3:1101:3300:94972#TGACCA/1
ACACTGGGACTGAGATACGGCCAGACTCTACGGAGGCGAGCAGTGGGGACTTTTGCCCAATGGGCGAAAAGCTTGACGCAGCAACGCCGCT
GATCGAAGAAGGACTTAGGTTCTGTAAGACTGTCTATAAGGGAAGAATAGCCAGTATTTAACACAATATTGGTCTGACGGTAC
>SOUFRE_0091:3:1104:15536:22926#TGACCA/1
TGCCCTTAGTACGGAATAGCCATTGAAACAGTATTAAATACCGTATACGCCCAAGGGGAAAAGTTTATCGCTAAAGGATCGGCCCGCT
TAGATTAGGTAGTTGGTAGGGTAATGGCTTACCAAGCCGACACTATAGCTGGTTTGAGAGGATGATCAGCAACTGGGACTG
>SOUFRE_0091:3:2107:7818:190958#TGACCA/1_rc
GTCAGCTCGTGTCTGAGATGTTGGGTTAAGTCCCACAACGAGCGCAACCCTACTTTTAGTTGCCACCATTAGTTGGGCACTTTAAAAGAA
CTGCCAGTGATAAGCTGGAGGAAGTGGGATGACGTCAAGTCTCATGGCCCTTAGTGTGGGCTACACACAGTGTCTACA
>SOUFRE_0091:3:1203:19509:71185#TGACCA/1_rc
ATAGAGGAAAGCAGAATTTCTAGTGTAGAGGTGAAATTCGTAGATATTAGAAAGAAATACCAATTGCGAAGGCAGCTTTCTGGATCAATACTGA
CACTGAGGAACGAAAGCATGGTAGCGAAGGATTAGATACCCTCGTAGTCCATGCCGTAACGATGTGTGTTAG
>SOUFRE_0091:3:2204:9225:53223#TGACCA/1_rc
GAATAAGCACCGCTAATTCGCTGCCAGCAGCCGCGTAATACGGAAGGTGCAAGCGTTAATCGGAATTACTGGGCGTAAAGCCGCGCTAGGT
GGTTTTGTAAGTTGAGTGTAAAGCTGGGCTAACCTCAACCTGATGAACCTGCATCAAA
>SOUFRE_0091:3:1304:11640:183863#TGACCA/1
GCTGGGCACTTAGAAAACTGCCGGTATAAGCTGGAGGAAGCGGGGATGACGTCAAGTCTCATGGCCCTTACGGTCTGGGCTACACAG
TGCTACAATGGTGGTACAGAGGGCAGCGATATCGAAGATAAAGCTAACTCCATAAAGCCATCTCAGTTCGGATTGGACTCT
>SOUFRE_0091:3:2308:13962:24638#TGACCA/1_rc
AGTGTAGAGGTGAAATTCGTAGATATTGGAAAGAAATACAGAAGCGAAGGCAGCTAACTAGGCCATTTTGACGCTGAGATACGAAAGCGTGG
GTAGCAACAGGATTAGATACCTGGTAGTCCACGCAAGTAAACGATGAGTGTAGTCTGCTGGGACATTA
>SOUFRE_0091:3:1108:20642:134730#TGACCA/1
GAATCAGCATGTCCGGTGAATACGTTGTAATACACACCGCCCGTACACCATGGAAGTGGATTGCACCAAGTATAGATAGTCTAA
CCTTCGGGAGGGCGTTTACCACGGTGTGCTTATGACTGGGGT
>SOUFRE_0091:3:1304:12009:32071#TGACCA/1_rc
ACGATCCATAGCTGGTCTGAGAGGATGATCAGCCACTGGGACTGAGACAGCCGAGACTCTACGGGAGGCAGCAGTAGGGAATTTGGA
CAATGGGGGCAACCTTGATCCAGCATGCCGCTGTGTGAAGAAGGCCCTAGGGTTGTAAGACACTTTCAGTAGGGAGGAAG
>SOUFRE_0091:3:1107:5254:143756#TGACCA/1
GTACTTACAATGGGATGCAAGAGGGCACTCTTAGTAACTCCCTAAAATGCACCTCAGTTCCGGATTGCACTCTGTAACCTCGATGATGAAGC
TGGAAATGCTAGTAAATCCGGGAACAGCGCCGGCGGTGAATACGTTCCCGGCTTGTCCACACCGCCCGTACACCATGGAAG
>SOUFRE_0091:3:2204:14834:84421#TGACCA/1
ACAAGTAGTGGGACGGTGGTTAATTCAAAGATACGGCAGAACCTTACCAACACTTGACATGTTCTGCGCACTTAAGAGATTAGATTT
TCGGTTCGGCCGAGCAAAACAGAGTGCATGCTGTCAGCTGTGCTGAGATGTTGGGTTAAGTCCCG
>SOUFRE_0091:3:1103:4218:5107#TGACCA/1
GTCCGACGGCCCTTCAAAGTCTGCTGTTAAAGCTGGAGCTAACTCATATGCGAGTGGAAACTGTTGGGCTTGGTGGTGGGGCA
GAGGAATTCCCGGTGATCGGTTGAAATGCGTATGCGGAAGAACCAGTGGCGAAGGCGCTCTGCTGGGCCATCACTGACGCTCAT
>SOUFRE_0091:3:2302:18191:165159#TGACCA/1
CCAGCAGCCGCGTAATACGGGAGTGGCAAGCTTATCCGGAATTTGGGCGTAAAGCGTCCGCGAGGCGCCCTTCAAGTCTGCTGTTAAAA
CGTGAGGCTTAACTCCATCATGGCAGTGGAAACTGTTGGGCTTGGTGTGGTAGGGGCAGAGGAATCCCGGTGATGCGGTGA
>SOUFRE_0091:3:2102:1868:98547#TGACCA/1_rc
TAAAGGCTTACCAAGGCTTCGATCAGTGTGCTGAGAGGATGATCAGCCACTGGGACTGAGACAGGCCAGACTCTACGGGAGGCA
CGACTGGGGAATTTCCGCAATGGGCGAAAGCCTGACGGAGCAACCGCCGCTGAGGAGCGAAGGCCCTTGGGCTGTAACCTCTT
>SOUFRE_0091:3:2107:20957:187106#TGACCA/1_rc
TGGTGGGTAATGGCTACCAAGACTGTGATCAATAGCTGATTTGAGAGGATGATCAGCCACTGGGACTGAGACAGGCCAAACTCTAC
GGGAGGCAGCAGTGGGGAATCTTGCACAATGGAGGAACTCTGATGACAGCGATGTGCGTGTGAGTGAAGAAGGCCTT
>SOUFRE_0091:3:1106:6610:58242#TGACCA/1
CGCTGAGGGACGAAGGCCTTGGGCTGTAACCTCTTTTCTCAAAGAAAGATATGACGCTACTTGGAGAAATAGCCACGGCTAATCCGT
GCCAGCAGCCGCGTAAATACGGAGTGGCAAGCTTATCCGGAATTTGGGCGTAAAGCGTCCGCGAGGCGCCCTTCAAGTCTGCTGTTAA
>SOUFRE_0091:3:2103:19422:74896#TGACCA/1
TGCAACTCGCTACGTGAAGTGAAGAATCGCTAGTAACTCGCAGGTGAGCATACTGCGGTGAATACGTTCCCGGCGCTTGTACACACCGCCGCT
ACACCATGGAAGTTGGCCG
>SOUFRE_0091:3:1107:13801:152822#TGACCA/1_rc
TGAGAACTAGCCGTTGGGAGATTTAACTGTTCAAGTGGCACAGTAAACGATTAAGTTCTCCGCTGGGGAGTACGGCGCAAGGCTAAAAT
CAAATGAATTGACGGGGGCCGCAAGTGGTGGAGCATGTGGTTAATTCGATGCAACGGAAGAACCTTACCAACCTTGA
>SOUFRE_0091:3:1207:17720:103115#TGACCA/1_rc
TGTAGAGGTGAAATTCGTAGATATTAGGAAGAATCAGAGGCGAAGGCGGCTCACTGTCGATGACTGACGCTGAGGTGCGAAAGCGTGGG
AGCAAAACAGGATTAGATACCTGGTAGTCCACGCGTAAACGATGGAAGTAGTGTGGACGGTTACTGTTCACTG
```

Sample Variance

#6 points

$$\sigma^2 = \frac{\sum_{i=1}^n (x_i - \bar{x})^2}{n-1}$$

Variance

Sigma means sum

$$(x_1 - \bar{x})^2 + (x_2 - \bar{x})^2 + \dots + (x_n - \bar{x})^2$$

↑ degrees of freedom

