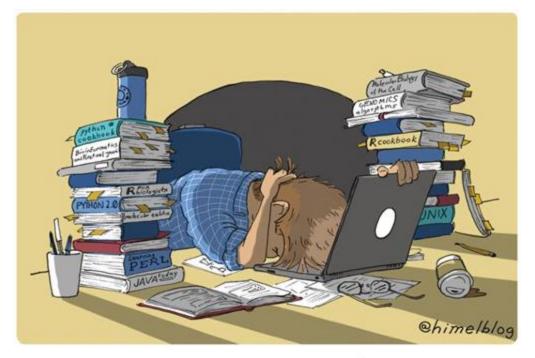
Reading, navigating, and plotting Cancer Research Institute on your own



Thoughts and challenges, troubleshooting, and working demo



"Maybe I should have stuck with Excel."

Ed Himelblau





How should the data be loaded?

- What type of data are we looking at?
 - What is the size of the dataset?
 - What are the variables?
 - Is the data complete? Why is data missing?
- What are the question(s) I'm trying to address?



How should the data be loaded?

- 1. Look at the file extension → Do you recognize it?
 - → No? Google "what is a <u>txt</u> file extension"
 Other extensions: csv, tsv, xls, Rda, Rds, h5, mtx



How should the data be loaded?

- 1. Look at the file extension → Do you recognize it?
 - → No? Google "what is a <u>txt</u> file extension"
 Other extensions: csv, tsv, xls, Rda, Rds, h5, mtx
- 1. Is a package required to look at this data?
 - Large txt or tsv files: data.table package → fread()
 - Excel files: readxl package → read_excel()
 - Google → What package can be used to read an <u>Rda</u> file in R? Is the package installed?



Reading in your data

```
library(data.table)
variant_file <- "/cloud/project/data/single_cell_rna/cancer_cell_id/mcb6c-exome-somatic.variants.annotated.clean.filtered.tsv"
print(variant_file)
read_variants <- fread(variant_file)</pre>
```

Make sure you include the full path of your file.

```
head(variant_file)
summary(variant_file)
nrow(variant_file)
ncol(variant_file)
```

Does the data look the way you expect it to?

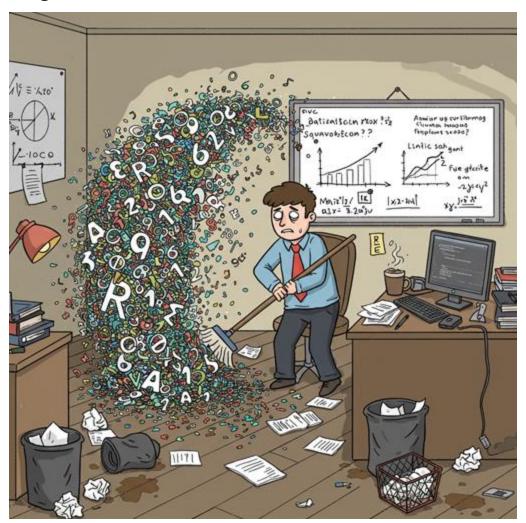


Is your data ready for analysis?



Is your data ready for analysis?

... Probably not



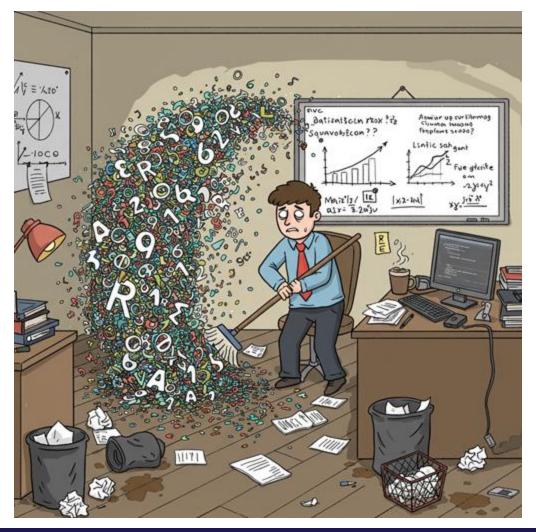


Is your data ready for analysis?

Data cleaning refers to the process of processing datasets to improve their quality for analysis and decision-making.

This may include:

- Removing or correcting missing information
- Identifying inconsistencies
- Performing batch effect correction





Missing data vs. sparse data

Why might data be missing?

- The file doesn't contain values for a given column in a subset of rows
- Computation/mathematical changes resulted in really large or really small numbers that can't be represented in memory
- You tried to add a column incorrectly
- Incomplete data file
- Technical reasons

Sparse data

Should there be a lot of zero values?



Missing Data

- NA ("Not Available" / "Not Applicable")
 - Generic missing-value marker in any vector type
- NaN ("Not a Number")
 - Result of undefined numeric operations (e.g. 0/0)
- NULL
 - Absence of a value or empty object (length 0)
- Inf / –Inf
 - \circ Infinite results (e.g. 1/0, -1/0)
- Character NA
 - Written as NA_character_ or as.character(NA) in character vectors







Detection functions

- \circ is.na(x) \rightarrow catches both NA and NaN
- \circ is.nan(x) \rightarrow isolates NaN only
- \circ is.null(x) \rightarrow tests for NULL objects
- \circ is.infinite(x) \rightarrow flags Inf/-Inf

Common handling strategies

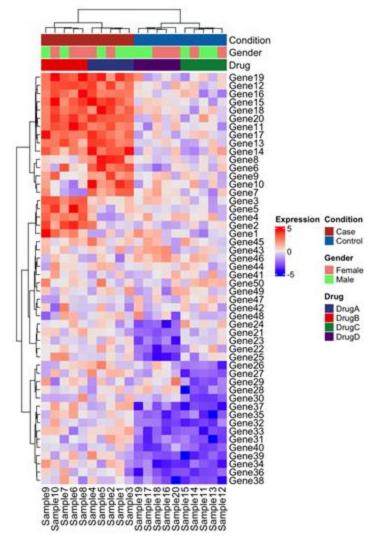
- o na.omit() / na.exclude() → drop rows with any NA
- o na.rm = TRUE → ignore NA in many base-R functions
- tidyr::replace_na() or dplyr::coalesce() → fill or impute missing values



Live demo

Replicating analysis by working backwards









- 1. Start your first Rscript
- 2. Work through the blocks of code under the Course: Reading, navigating, and plotting data page
 - Review the details on how the code works in the Lecture slides for assistance
 - Put a post-it on your laptop if you get stuck, indicating for a TA to come up to you
 - Work through the blocks of code on this page, practicing in both your Rscript and the console
- 3. Take the next step
 - There are a list of Additional exercises at the bottom of the page for you to try on your own

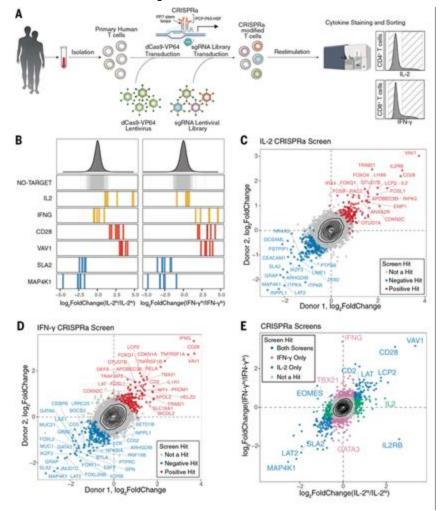


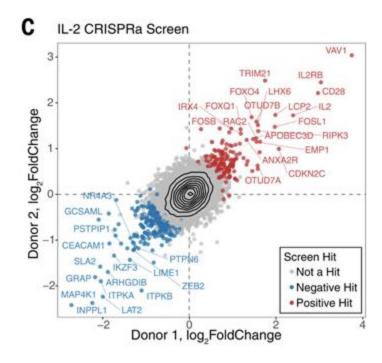
Goal: Have the foundation to take on your own dataset

Can you load one of your own datasets into R?



Live demo (if time)





Schmidt, et al. Science 2022