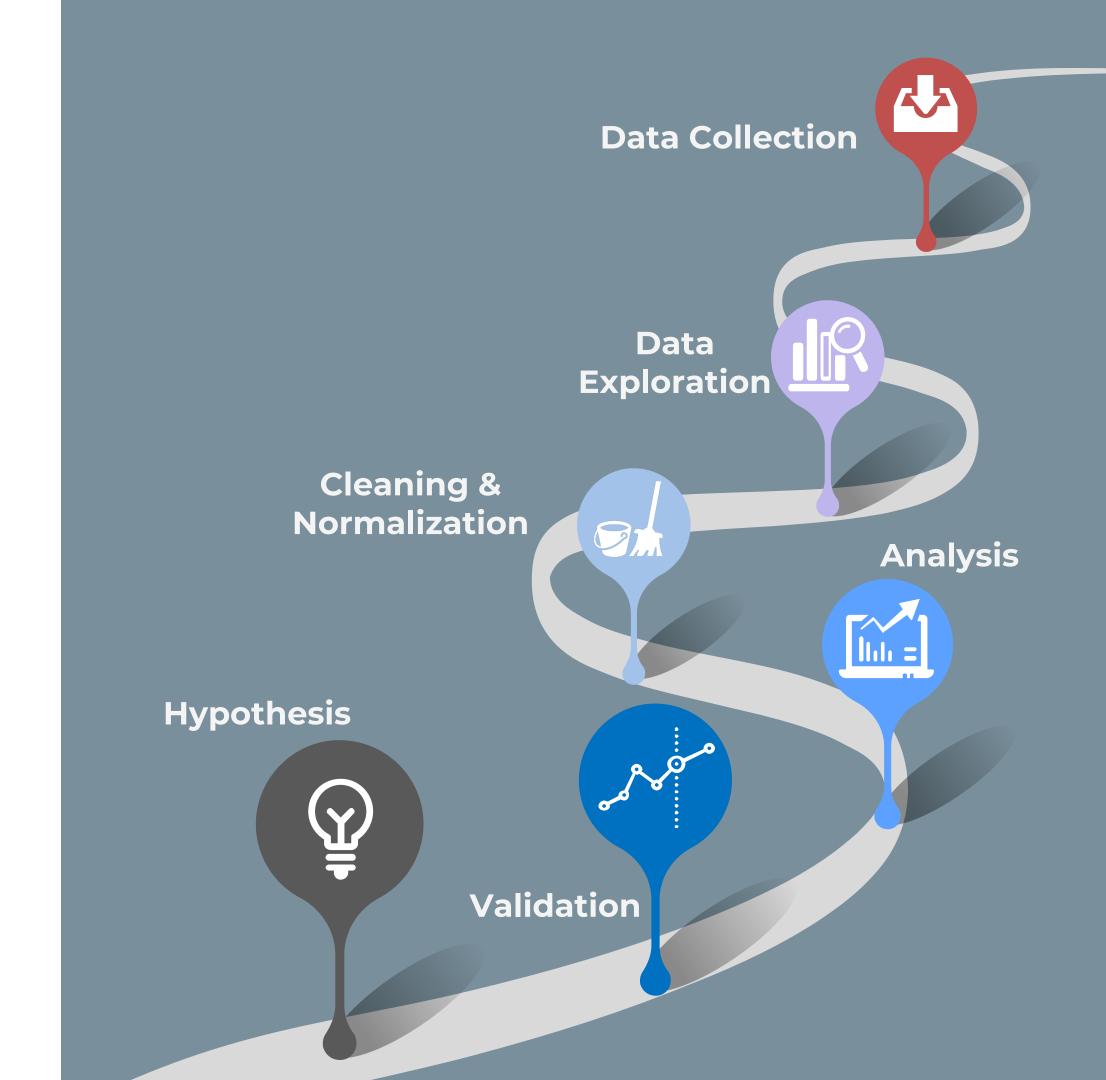
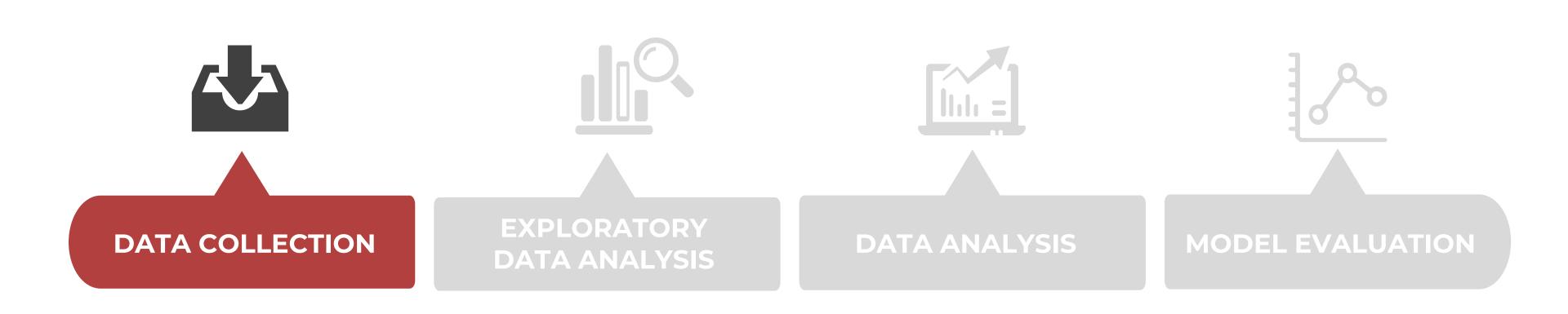
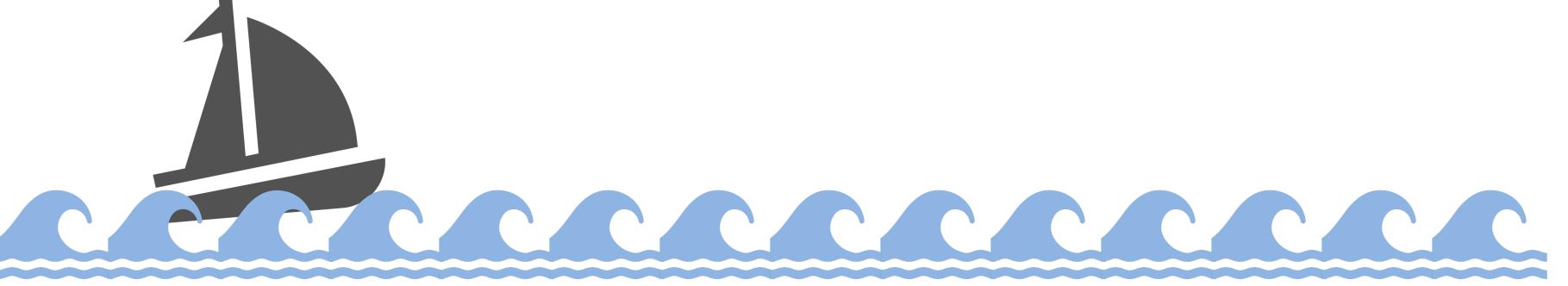
THE DATA'S JOURNEY





BEGINNING OUR JOURNEY

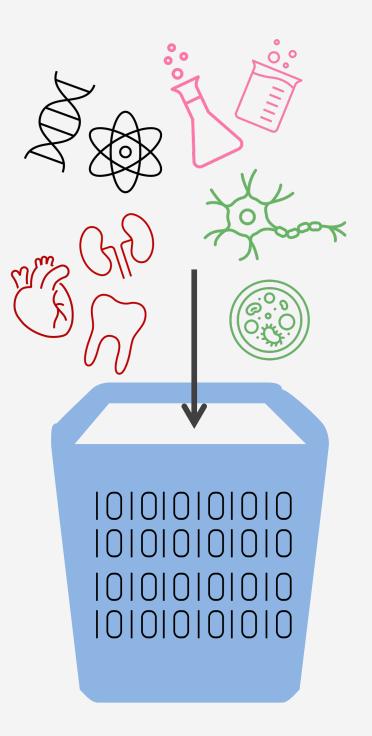






DATA COLLECTION & TERMINOLOGY

- Data points are **observations of reality**, made with the help of measuring devices or techniques which means:
 - Measurement uncertainty (technical variation)
 - Introduced biases
- Type and impact of of these may be known or unknown
- Datasets:
 - Observations, the properties collected, i.e. a cell, a patient sample
 - Variables (Features), i.e. blood pressure, smoker status, gene expression
 - Outcome variable(s) (dependent)
 - Explanatory variables (independent)?





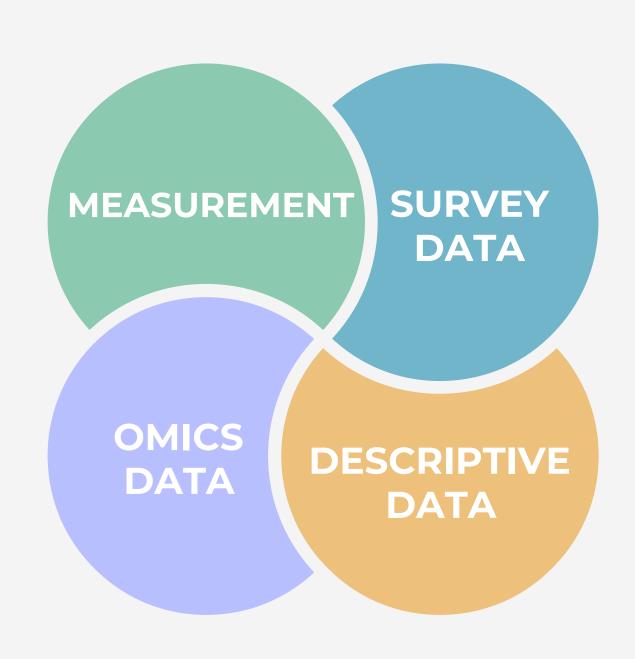
DATA TYPES IN HDS

Measurements

- i.e. height, weight, heart rate
- Represented as a number or a category
- Often low dimensional, but wearables are changing that

Omics Data

- i.e. genomics, proteomics
- (Ultra) high-dimensional
- Needs a lot of preprocessing



Survey Data

- Can be descriptive, numeric or categorical
- Relies on reporting, i.e. very prone to bias

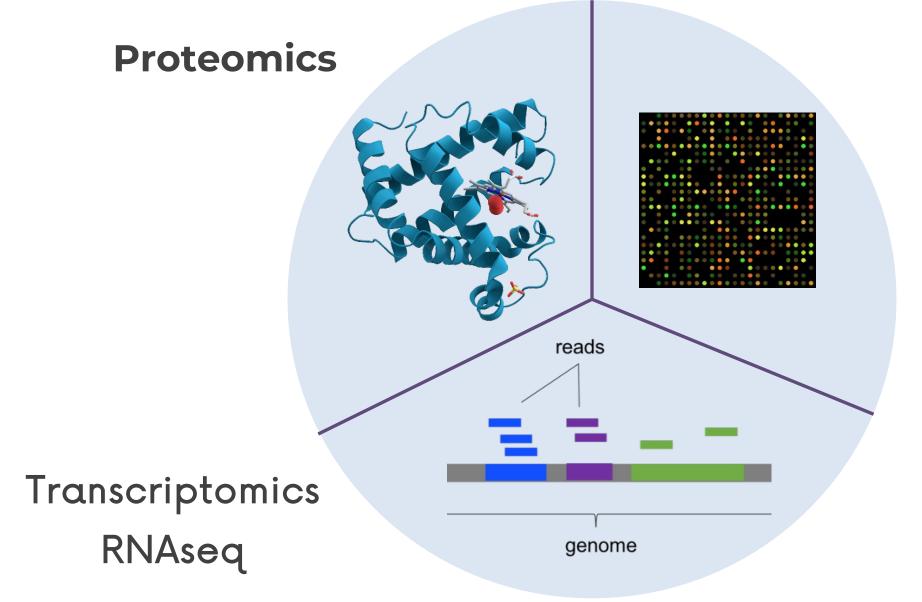
Descriptive Data

- i.e. patient journals, registry data
- Highly person-sensitive
- Not easily standardized



OMICS DATA

Omics approaches aim to study the entirety of an 'ome' (proteome, transcriptome, genome). Here we name the most commonly used types.



Whole genome seq
SNP calling



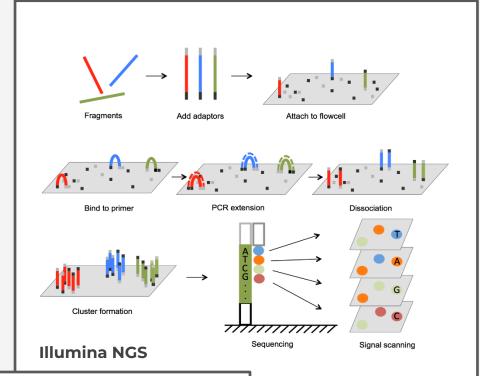
OMICS DATA

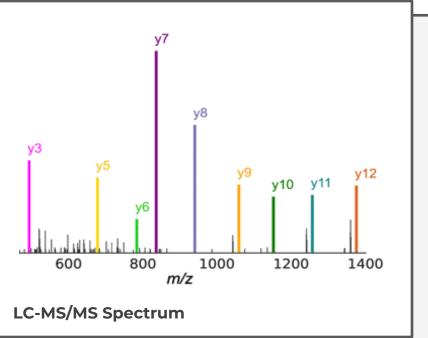
Omics data are produced with specialized lab protocols followed by:

- Next generation sequencing (DNA, RNA)
- Mass spectrometry technologies (proteins, lipids, sugars, metabolites)

Data characteristics:

- Extremely high-dimensional
- Expensive to generate
- Measurement is indirect
- Need for pre-processing
- Can be prone to high variability (therefore replicates)







BIOBANKS & REPOSITORIES

- Biobanks and repositories are great sources of both bio data and patient metadata
- Great resources in Denmark but access can be a cumbersome process, as these are (highly!) person-sensitive data



 Access has to be applied for through proper channels and compliance has to be ensured while working with these data.

HeaDS GDPR course for biomedical researchers!



REGISTRY DATA

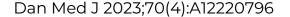
Most doctors, epidemiologists, ..., and statisticians work with registries

Data characteristics:

- Data are sensitive & hard to access
- There are many inconsistencies and errors
- Missing data are very common
- Needs a lot of clean-up and restructuring

The registries are governed by **Sundhedsdatastyrelsen** (The Danish Health Data Authority).







EXPERIMENTAL DESIGN

You have heard it before and you will hear it again, experimental design is important.

WHAT IS GOOD EXPERIMENTAL DESIGN?

- True normal controls
- Power calculations
- Randomization (data collection & laboratory)



Documentation



WHY IS IT SO IMPORTANT?

- You will be able to correctly answer your scientific question
- Saves you time/work
- Saves you money





CONFOUNDING

We often see non-randomized design, and sometimes total **confounding**.

Confounding: We cannot distinguish the separate effects of two different sources of variation.

There is **no way to correct confounding** when it is complete (this means the two variables 100% co-occur).

Hi I am Jake.

I have proteomics samples from people with diabetes and healthy controls. Can you help me analyze these data? They cost us 250.000 DKK to produce.



Hi, I am Thilde. I would love to! Where were the samples collected & stored?





Jake my friend, I have some bad news....





GROUP DISCUSSION 1.1

Consider the following scenario (1):

You would like to study whether smokers have an increased risk of heart disease. For this, you have collected some data from both men and women on whether they smoke and have heart disease.

When you examine the data you see that all of your smokers are men.

Can you answer your research question with this dataset?



GROUP DISCUSSION 1.2

Consider the following scenario (2):

You would like to study differences in gene expression between tumor and healthy tissue. Since you have a lot of samples you ask two lab techs to each process half.

What are potential biases and confounders in this set-up?



BEFORE THE ANALYSIS COMES...

- File formats in data science usually need to be a table (csv, xlsx, tab delimited, ...)
- Omics platforms return:
 - Fasta, fastq, SAM, GFF, MzML, ... huge files do not fit in DS analysis
- Registries are:
 - In formats more easily legible by humans, but still poorly structured for DS
- At the point where we do what is typically regarded as DS, the raw data has been translated into the quantities we want to measure and/or restructure.
- The steps we take before Data Science may be called: data management, set-up or wrangling



DATA MANAGEMENT

DOS

Write down steps (protocol & comments)

Record date of downloads

Consistency of file naming and folder structure

The raw data is untouched!

Consistency of data management

Don't store BIG DATA in your computer

Avoid XLSX spread sheets

No color coding

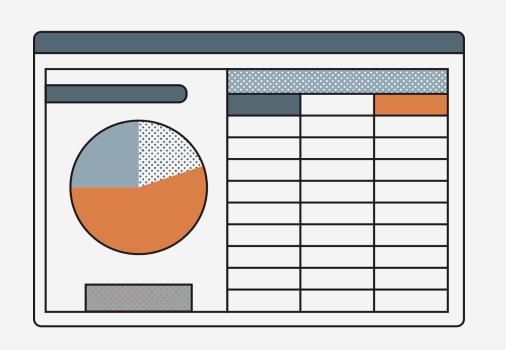
No spaces nor special characters in naming

Do not do things 'manually' if avoidable

DON'Ts



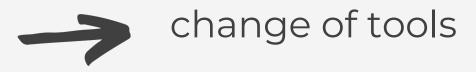
NEW TOOLS FOR DATA ANALYSIS







- Absolutely massive amount of data! Both in terms of number of observations as well as number of measured variables
- New types of data (i.e. omics, geolocation, wearables, ect)

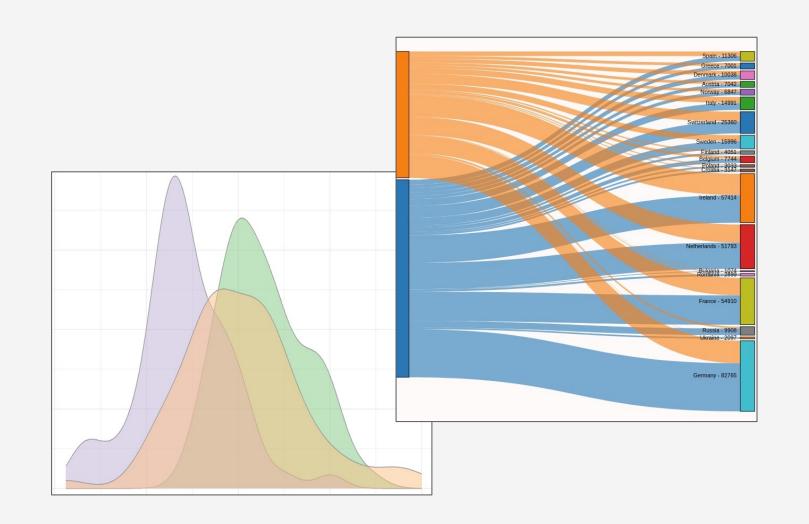


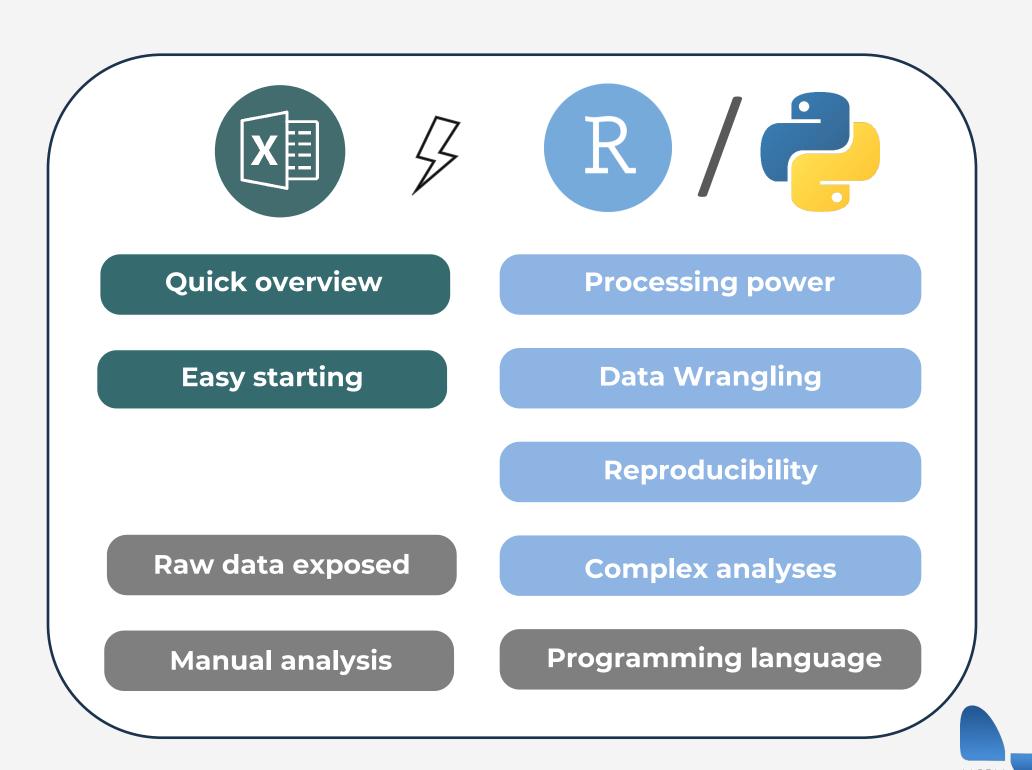




PROGRAMMING LANGUAGES

 Part of the new tools is the use of formal programming languages to analyze data instead of tools like Excel.







Q5: Which programming languages

have you had contact with?





THE DATA-HYPOTHESIS CYCLE

Hypothesis-driven:

- 'traditional' way of research
- formulate hypothesis
- design experiment to challenge
- hypothesis supported or rejected

Data-driven:

- Discover properties of data set
- Identify patterns and relationships
- Mostly possible with BIG datasets
- data-driven ≠ fishing!

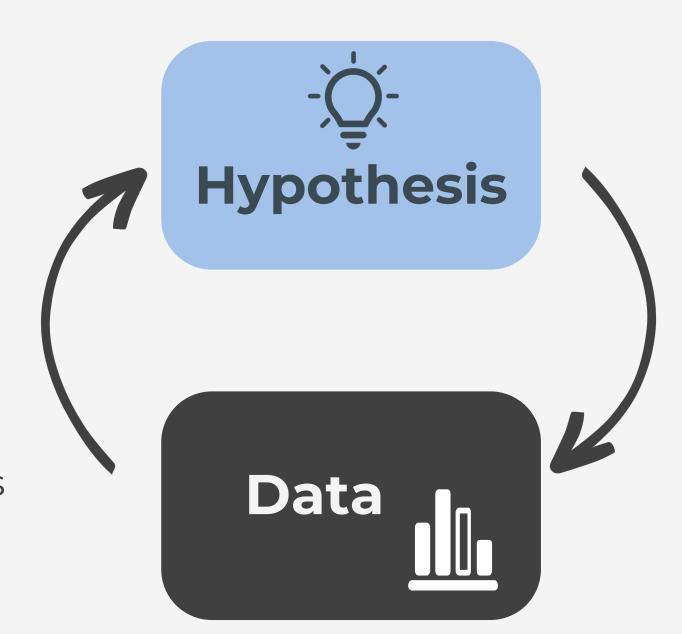




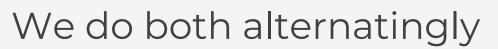
THE DATA-HYPOTHESIS CYCLE

High-dimensional data with many observations

- Discover complex patterns that humans cannot identify without algorithms & computer power
- Find new patterns from re-analysis with new (multiple) methods or combination of datasets



Gather new data to confirm or deny new hypothesis

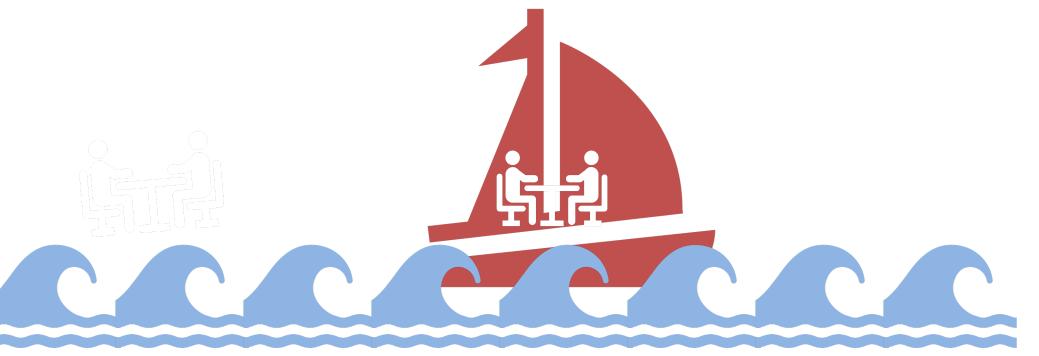




GROUP DISCUSSION 1.3

In your groups discuss:

- What **data types** do you and your collaborators currently work with and/or what are you interested in working with in the future?
- What considerations are there in terms of experimental design, data collection and/or data management/set-up?





BREAK

