

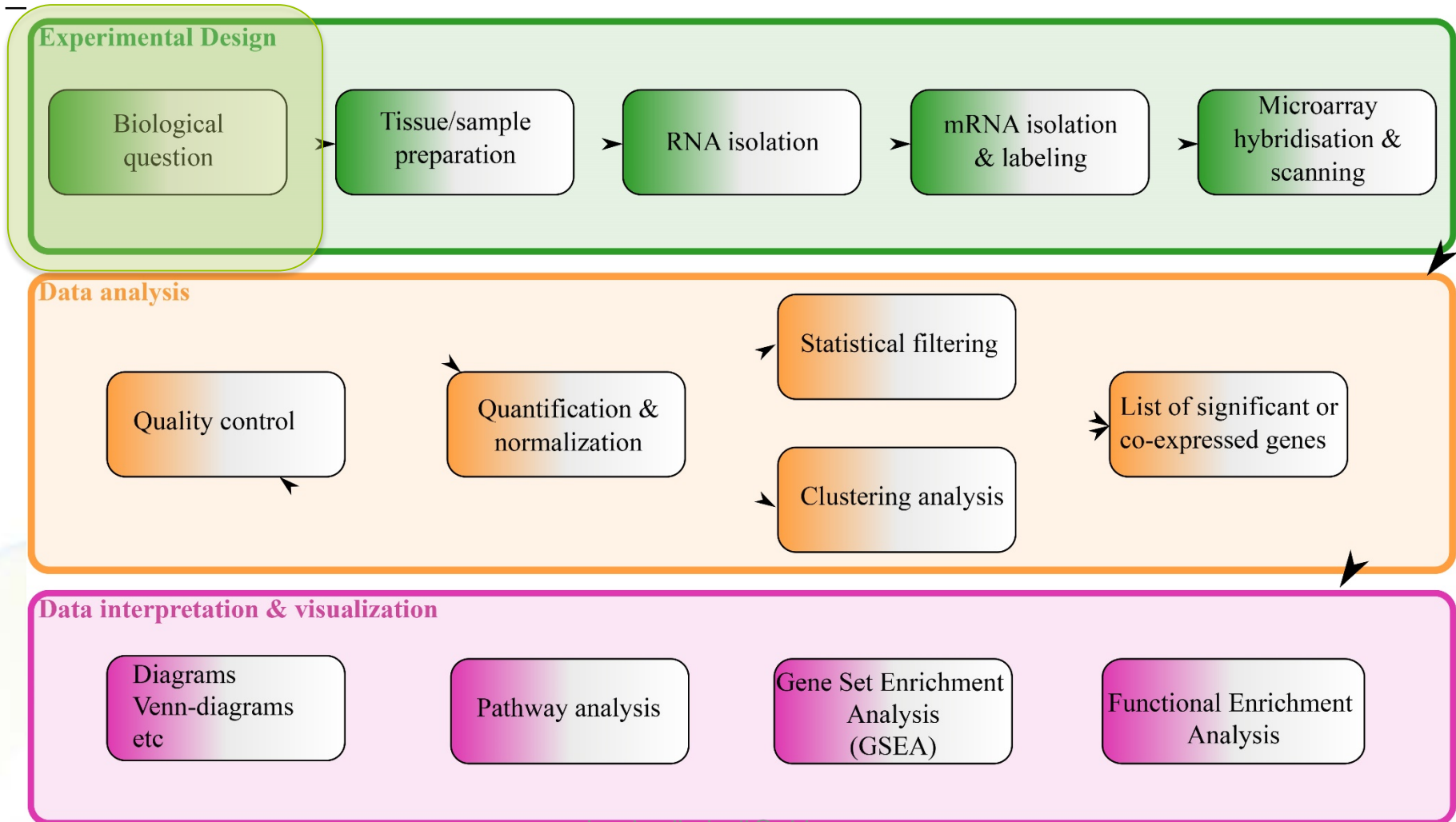
Towards different biological conditions - what to think about

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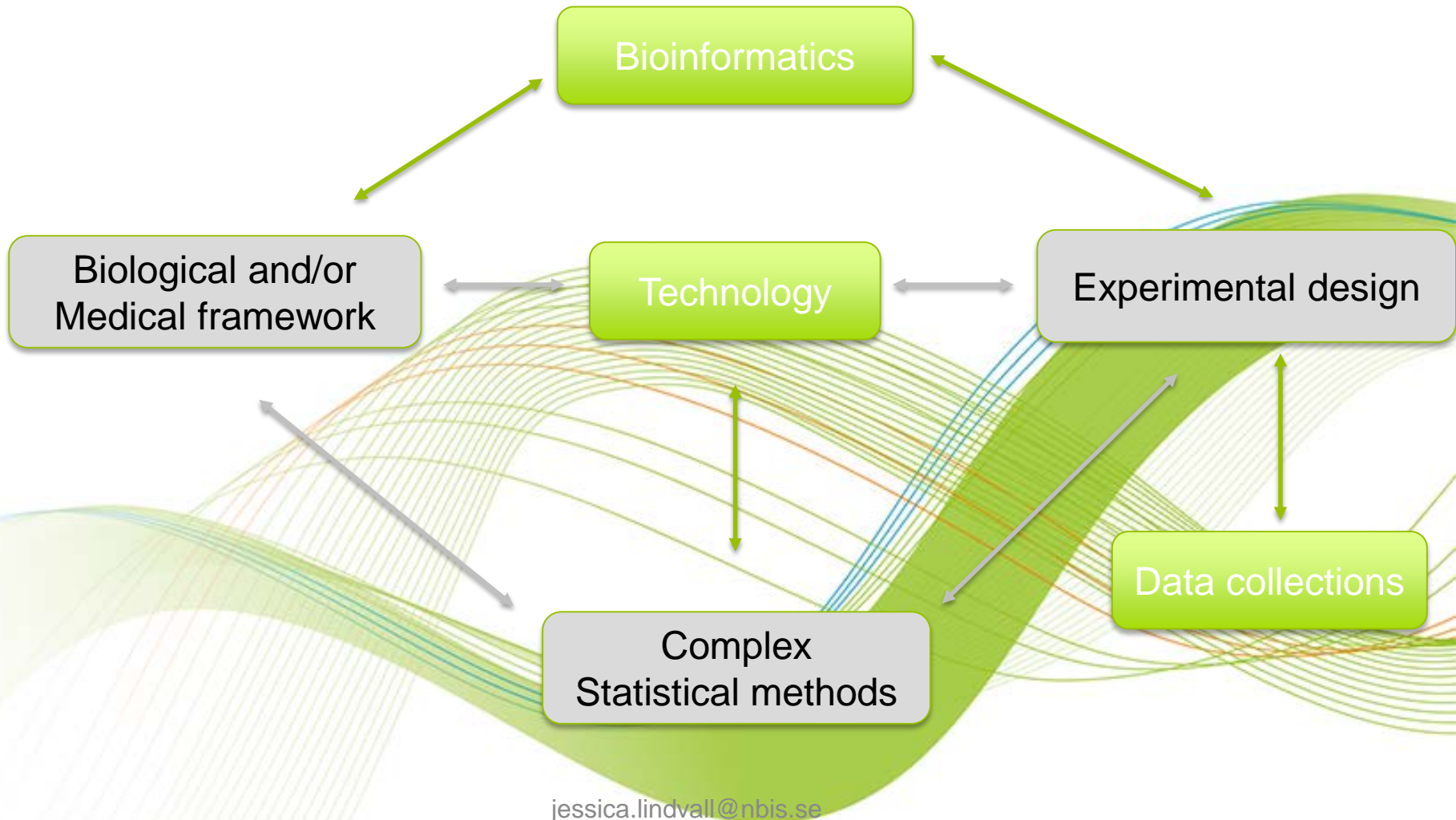
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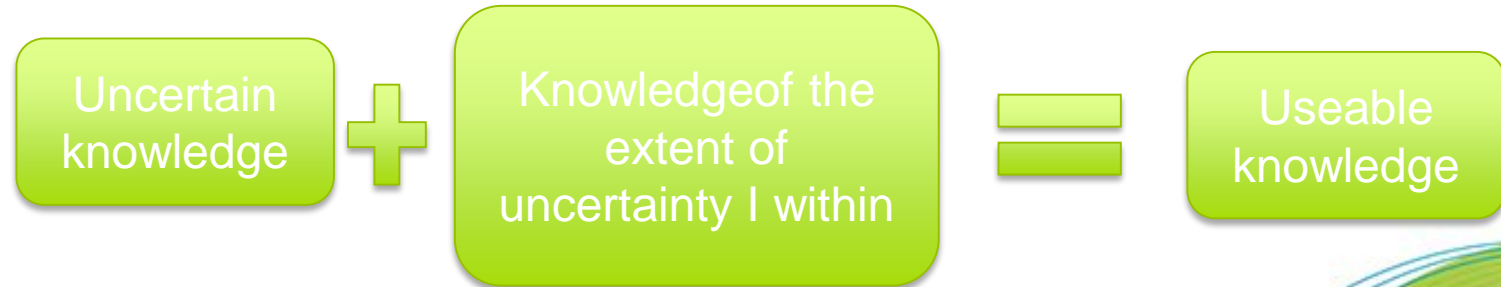


Microarray Data Analysis Workflow



It all goes together!





Measurement Model

$m = \neq + e$

m – measurement model

\neq - True, however UNKNOWN value



Decisions on the
Experimental
design influence the
measurement model

What is the *meaning* of e ?

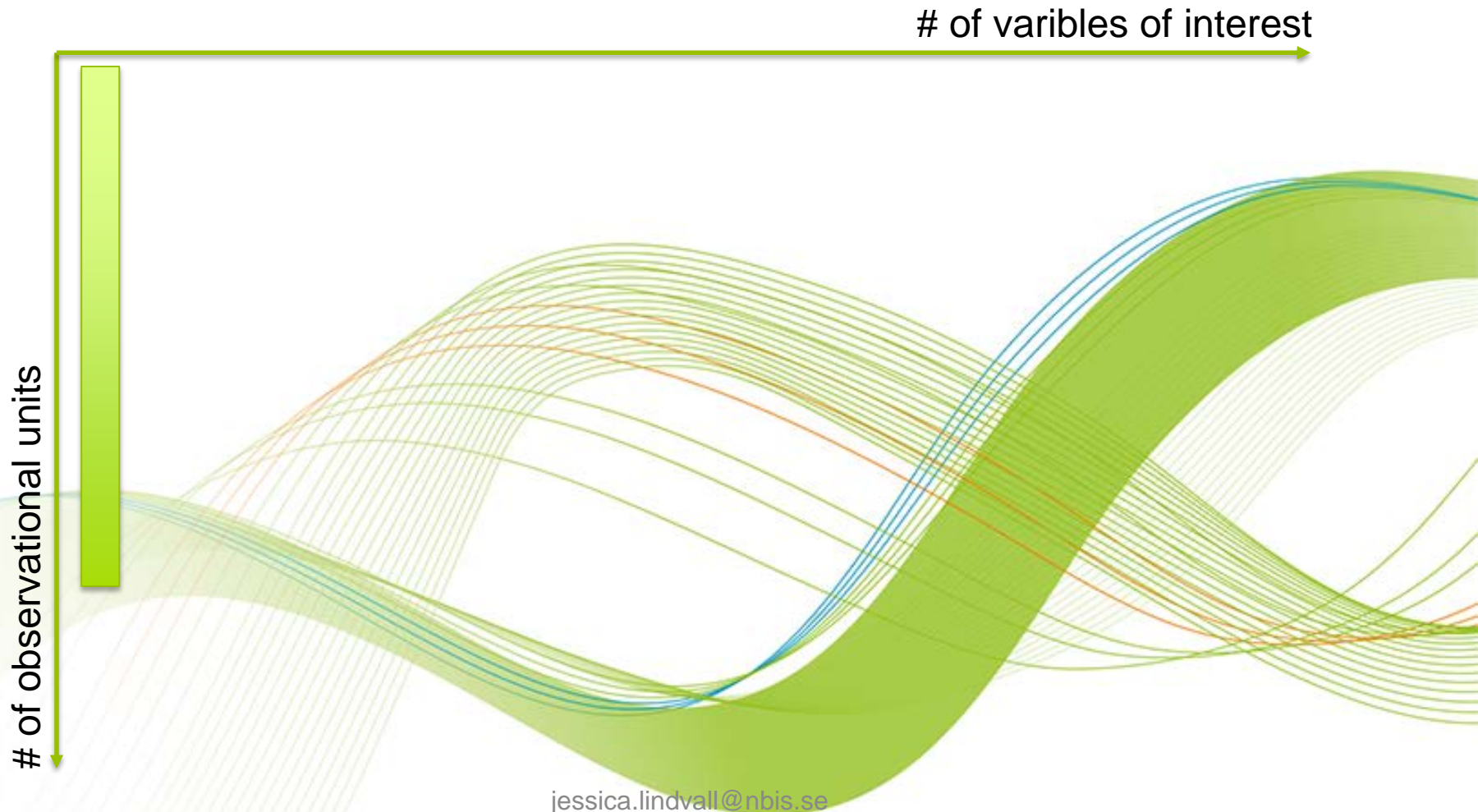
What is the *variance* of e ?

Dependence between e and \neq

Distribution of e and \neq



Information dilemma – too many or too few?





Information dilemma – too many or too few?

of variables of interest

of observational units

The experimental set-up – a general take!

- Scientists deal mostly with experiments of the following form:
 - A number of alternative conditions / treatments,
 - one of which is applied to each experimental unit,
 - an observation (or several observations) then being made on each unit.
- The objective is:
 - Separate out differences between the conditions / treatments from the uncontrolled variation that is assumed to be present,
 - Take appropriate statistical steps towards understanding the phenomena under investigation.

Experimental design – From a Biologist perspective

- *know the pitfalls and assumptions* of particular design,
- be able to identify the type of *model appropriate for the sampling design and type of data* that will be collected,
 - Species
 - Tissue/Blood
 - Primary cells
 - Cell lines
- be able to biologically *interpret the output* of analyses using the statistical model,
- be able to *design experiments optimally*, i.e. with the best possible use of our *limited time and resources*.



Experimental design – From a Statistician perspective

- Description and detection of the pattern MUST BE done in a rigorous way,
 - Detect gradients in space and time -> models explaining the pattern,
 - High confidence in estimations of the parameters in the statistical model
- Hypothesis testing of the design and model of the experiment is crucial.



Statisticians and Biologists – the common language

Statistician to the Biologist

- frame our questions in such a way as to get a sensible answer (statistical solidity),

Biologist to the Statistician

- be aware of biological considerations that may cause statistical problems
- understand the advice or analyses that we receive, and be able to translate that back into biology

Main requirements for an experimental setup

Once the *conditions / treatments*, *experimental units*, and the *nature of the observations* have been fixed, the main requirements are:

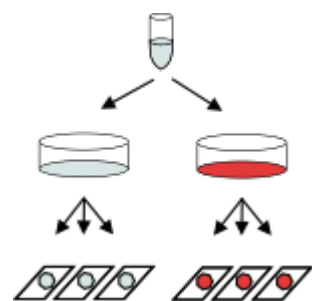
- Experimental units receiving different treatments should differ in no systematic way from one another – *Assumptions that certain sources of variation are absent or negligible should, as far as practical, be avoided*;
- Random errors of estimation should be suitably small, and this should be achieved with as few experimental units as possible; .
- The conclusions of the experiment should have a wide range of validity;
- The experiment should be simple in design and analysis;
- A proper statistical analysis of the results should be possible without making artificial assumptions.

Raise awareness – what to take into consideration

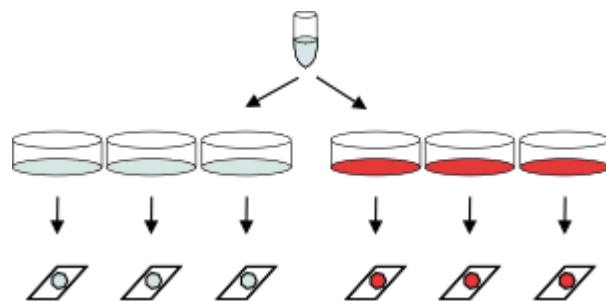
- General thoughts – always applicable!
- 3 different examples – different (biological) ways of thinking!
 - Established (immortalized) cell lines
 - Human primary cells
 - Animal models e.g. mice or flies



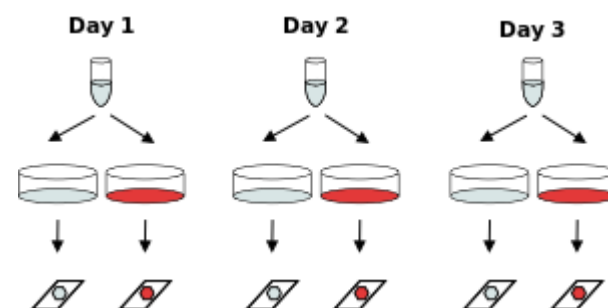
Aim for “true” Biological replicates



BAD



BETTER



GOOD ENOUGH

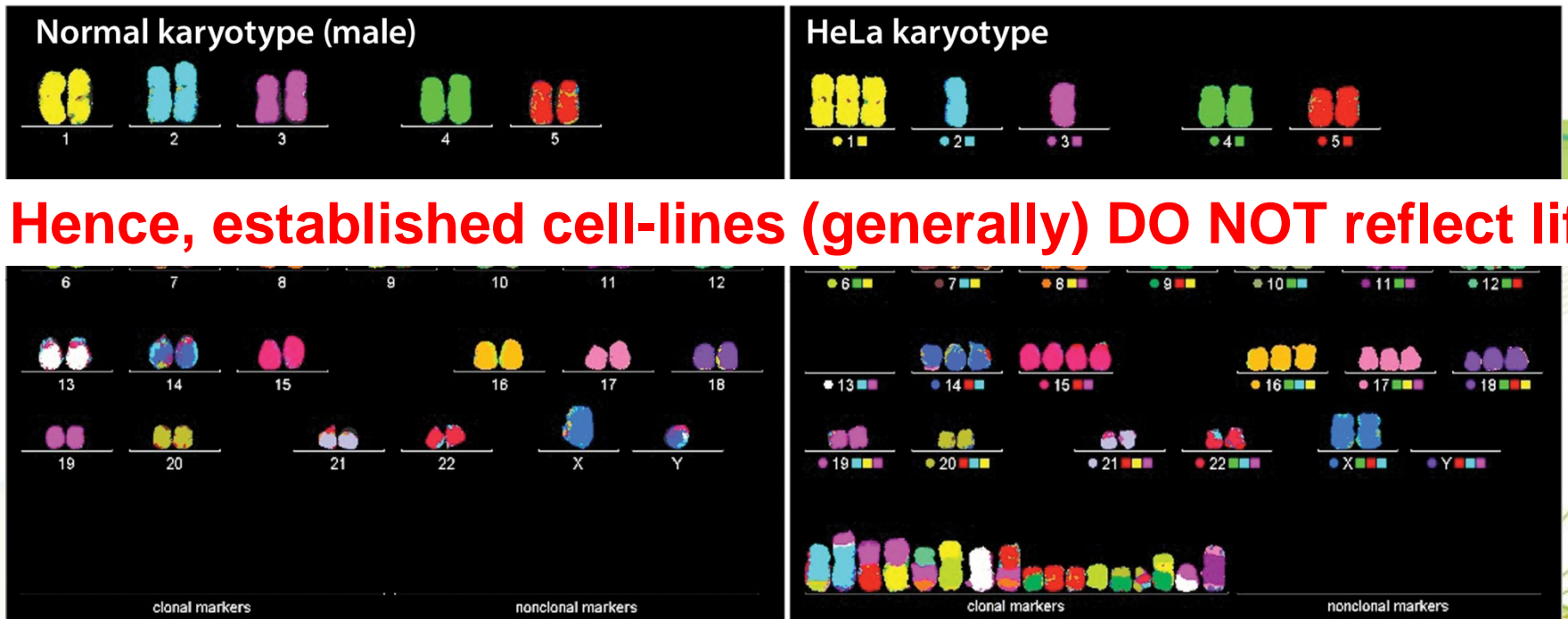
- The *ideal* design would have *biological replicates (i.e. cells from multiple people or animals)*

Established cell-lines

- HeLa cells
 - Transformed/immortalized cell line -> strongest clone “wins”!
 - Number of passages different from different labs -> strongest clone “wins”!
 - (Not the same “winning” clone in the different labs (or not even within the same lab maybe))

Established cell-lines

- HeLa cells



Hence, established cell-lines (generally) DO NOT reflect life

Do a Karyotype test before your experiment to be aware of what you have!

Human primary cells

- Potentially there is a mixture* of cells e.g. blood/PBMC
- Better if one can purify specific cell types e.g. B-cells from blood etc.
 - * *there are tools to handle this*
- “Cleaner” – no messy karyotypes- reflecting “true” biology
- Harder to collect (maybe) Need higher number of replicate samples per group – to mirror the true population

Animal models

- More controlled compared to Human primary cells/tissues
- Strain background *might* (IS) make a difference in outcome
- Less replicates are needed (due to more control) compared to human primary cells/tissues
- Number of animals might increase – Sufficient number of primary cells hard to obtain?!?



Biological data usually messy!

- Use statistics as a tool to clarify the biological question
- Design of the project important **BEFORE** starting the experiment
 - Starting material (factors influencing the data output)
 - Mixture of cell types
 - Transformed cell-lines (messy karyotypes)
 - Gender, Age...
 - Batch effects (experimental date, lab effects, technology)
 - Awareness of your data takes you a long way!

One last thing to keep in mind – The picture!



Questions?