

# AI LAB

1.3&4 – LET'S LOOK AT SOME DATA FOR REAL

FRANCESCA M. BUFFA



# LAB STRUCTURE

- INTRODUCTION
- THE DATA
- THE AI-LAB CHALLENGE – PART 1
- PART 1 - SHARING AND DISCUSSION

- UNSUPERVISED LEARNING EXAMPLES
- THE AI-LAB CHALLENGE – PART 2
- PART 2 - SHARING AND DISCUSSION

- SUPERVISED LEARNING EXAMPLES
- THE AI-LAB CHALLENGE – PART 3
- LARGE PROJECTS AND DATABASES

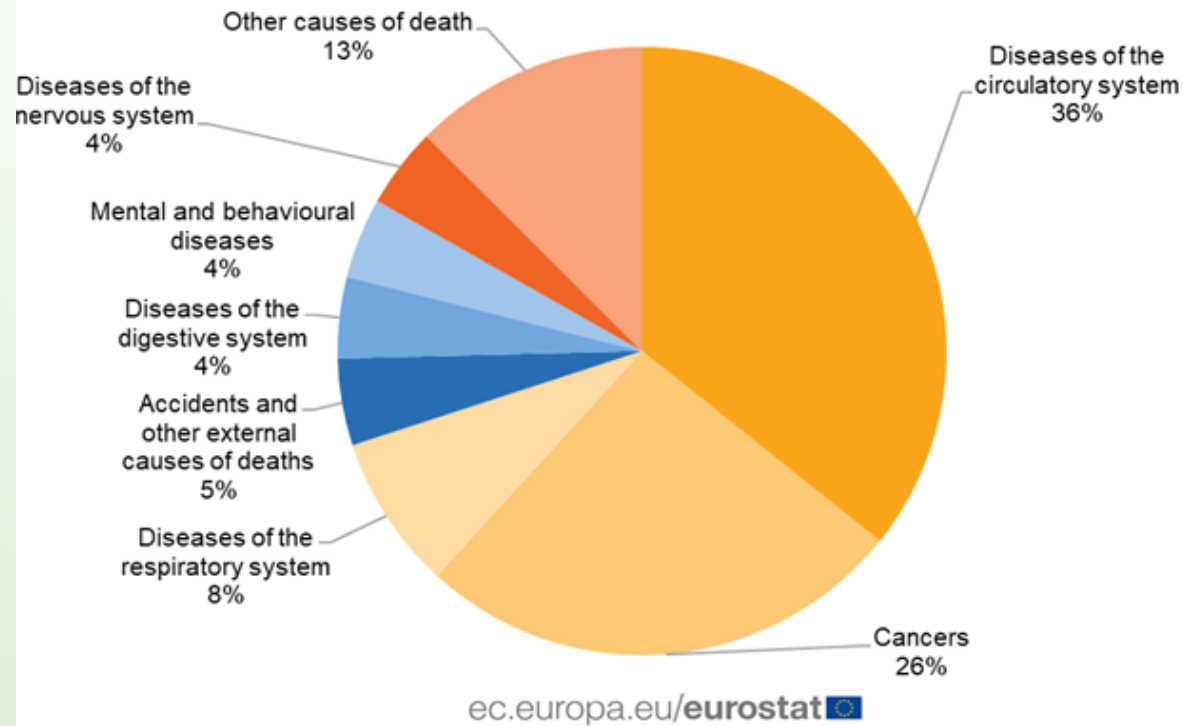
- THE AI-LAB CHALLENGE PARTS 1-3, SHARING AND DISCUSSION
- DATA INTERPRETATION
- DISCUSS AND PREPARE WORKSHOP PRESENTATIONS

# CHALLENGE - PART 1

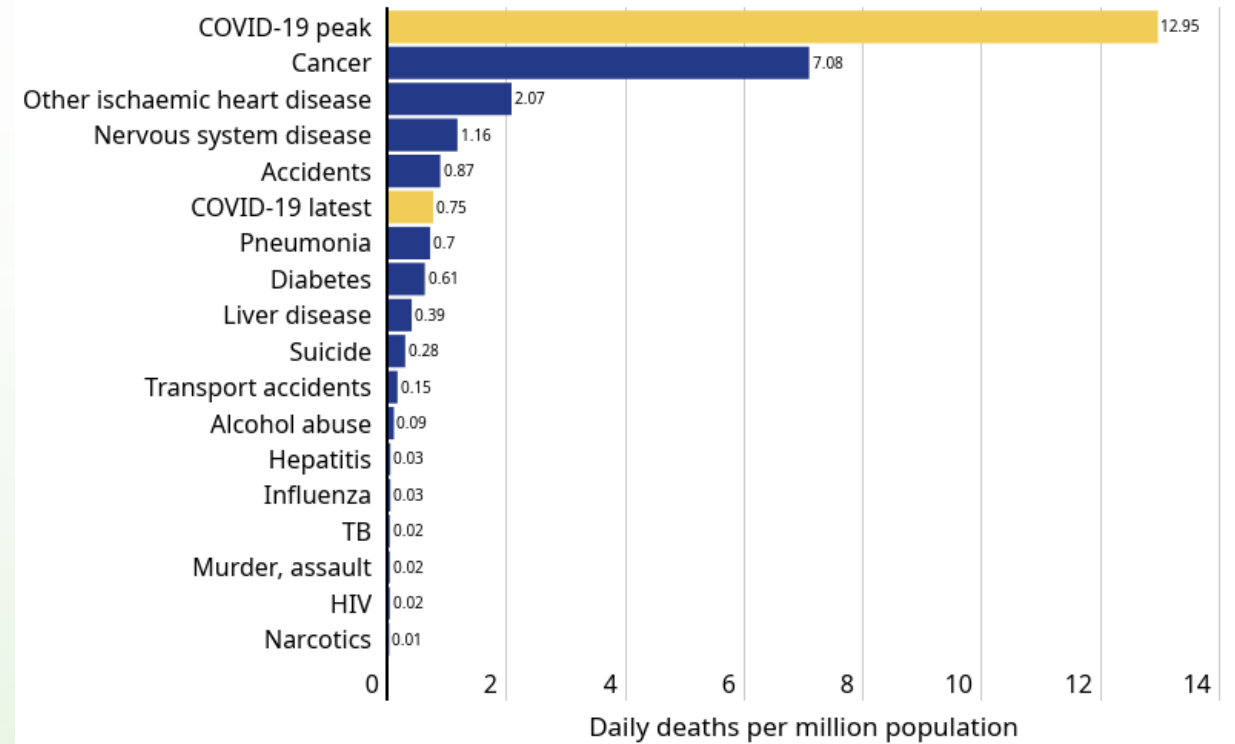
- DIVIDE INTO GROUPS, 4-5 PEOPLE PER GROUP IF POSSIBLE
- DOWNLOAD 3 GENE EXPRESSION MATRIXES FROM LINK-D
- THE DATA YOU WILL DOWNLOAD ARE CONFIDENTIAL AND FOR USE IN THIS COURSE ONLY
- LOOK AT THE DATA DISTRIBUTION (PLOT, CALCULATE DESCRIPTIVE STATISTICS, ETC)
- LOOK AT THE DATA STRUCTURE (PCA, CLUSTERING)
- COMPARE THE 3 DATASETS
- WRITE REPORTS WITH YOUR INITIAL COMPARISON OF THE 3 DATASETS,
- INCLUDE THE CODE YOU USED TO GENERATE THE REPORTS
- UPLOAD YOUR REPORTS TO LINK-U

# SOME BACKGROUND: CANCER

Causes of death in the EU by type, 2016  
(as % of all deaths)



Latest figures show the pandemic is still killing more than many causes of death

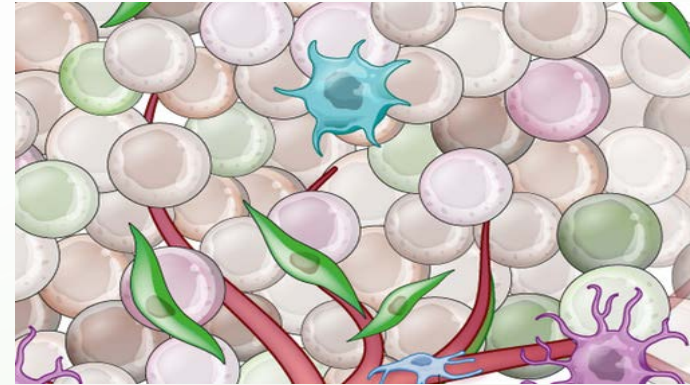


Source: Eurostat, Johns Hopkins University.

# THE CHALLENGE

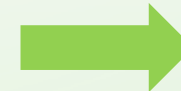
Cancer is a complex and heterogeneous disease

1. genetics
2. cell-cell signalling
3. microenvironment



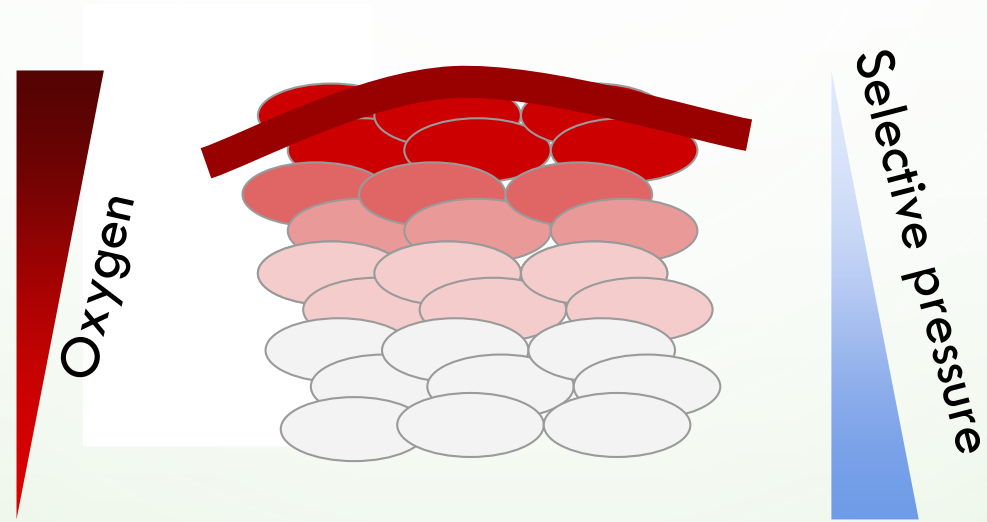
Many FDA approved drugs available

<i>Cetuximab</i>	<i>EGFR</i>
<i>Bevacizumab</i>	<i>VEGF</i>
<i>Olaparib</i>	<i>PARP</i>
<i>Enzalutamide</i>	<i>AR</i>
<i>Pictilisib</i>	<i>PI3K</i>



Therapeutic resistance

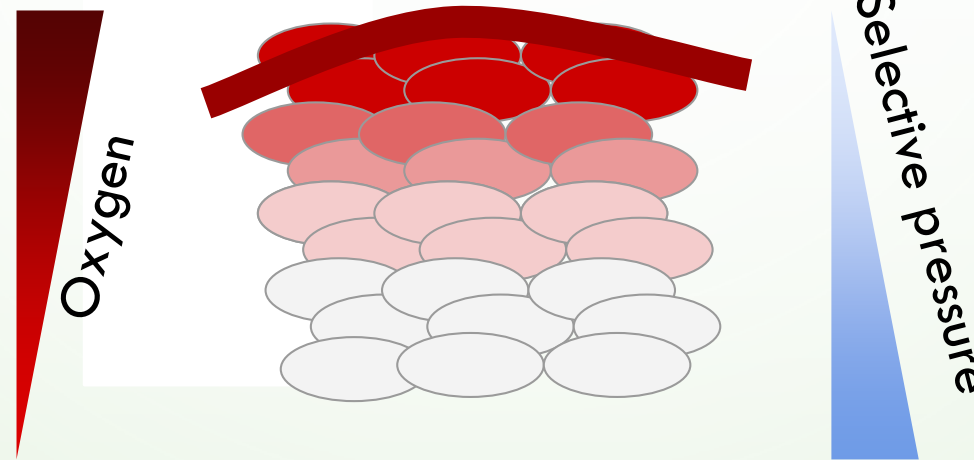
# A hypoxic tumour microenvironment





# A hypoxic tumour microenvironment

Hypoxia key trait of  
the tumour  
microenvironment  
[Genome Med \(2011\)](#)



Key methodology  
to identify hypoxic tumours  
[Cancer Cell \(2013\)](#)

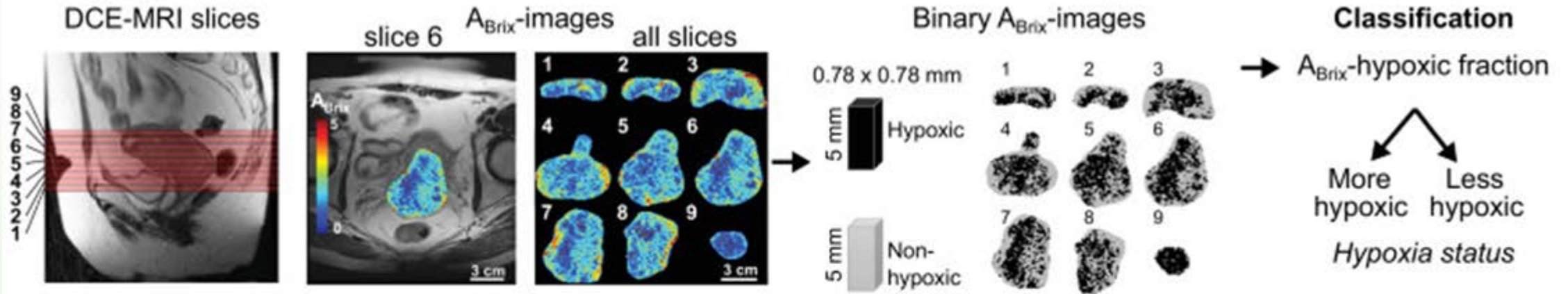
Hypoxia, poor prognosis  
and treatment resistance  
[JNCI \(2011\)](#)

Hypoxia, genomic instability  
and selective pressure  
[Genome Biology \(2016\)](#)

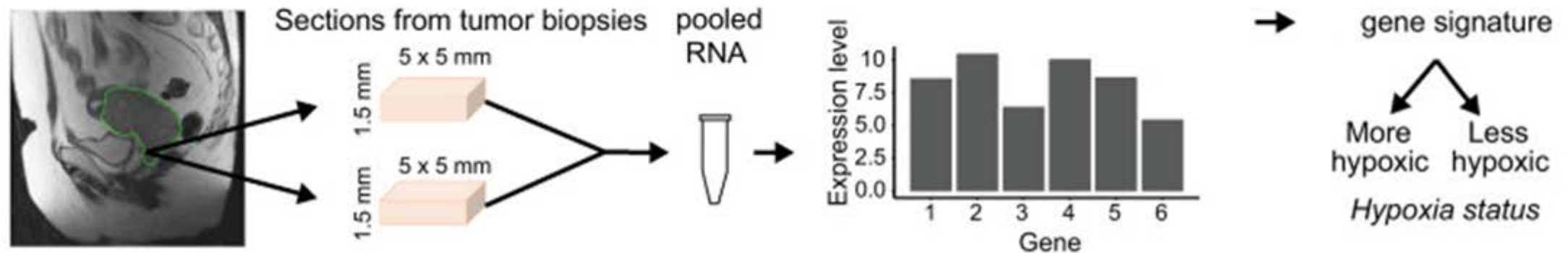
*Can we identify hypoxic cells?*

# HOW DO WE MEASURE HYPOXIA IN TUMOURS?

## Imaging-based biomarker

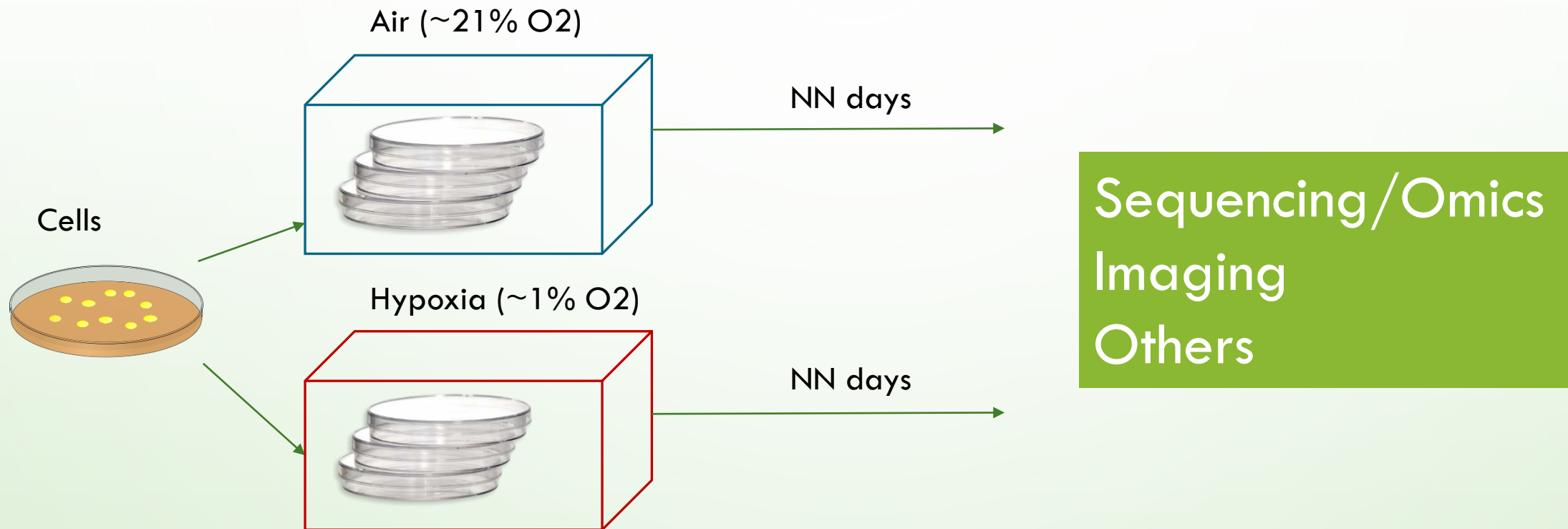


## Gene-based biomarker

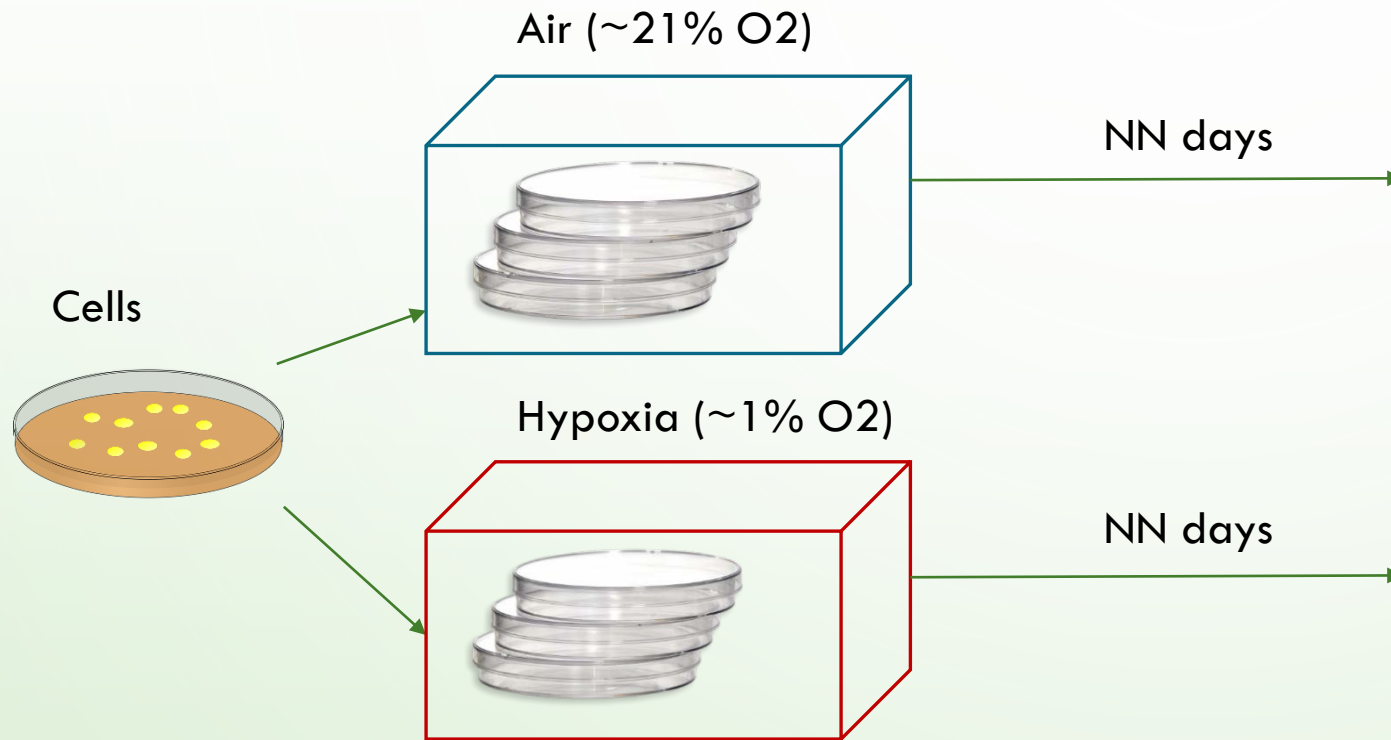




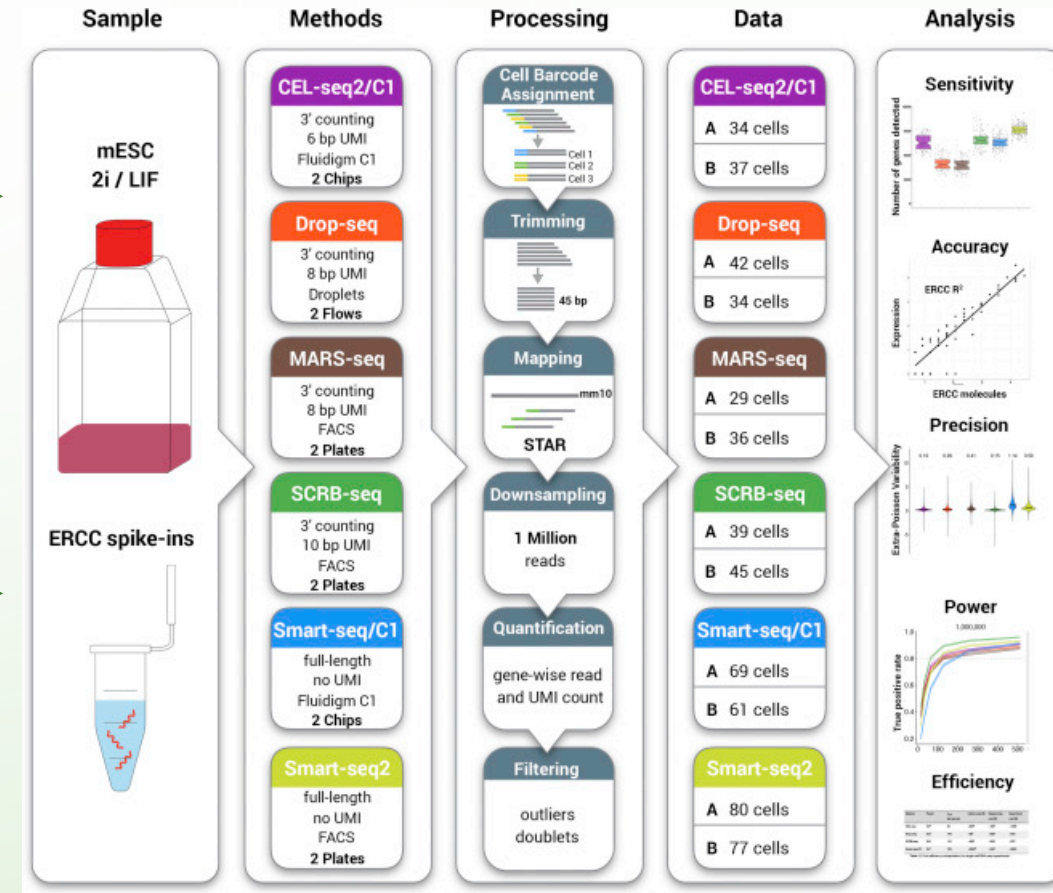
# DERIVE GENE BASED BIOMARKER IN THE LAB



# THE EXPERIMENTS WE WILL ANALYSE

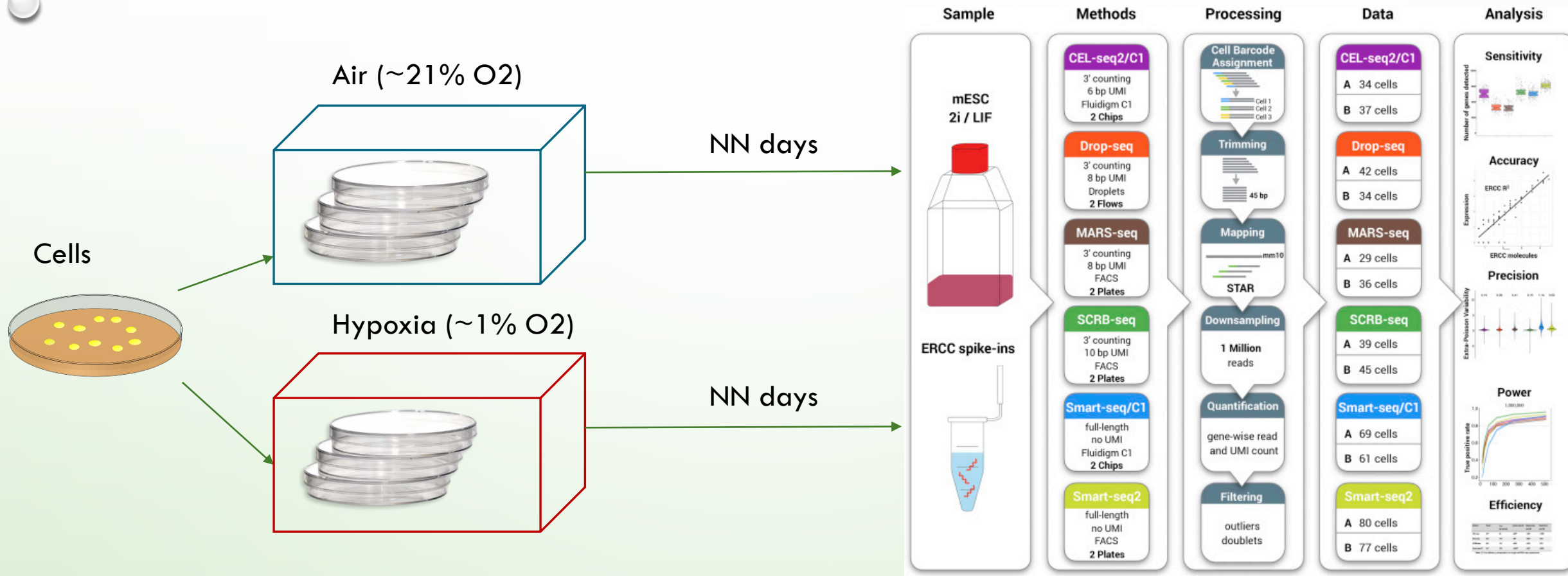


## Single-cell RNA sequencing



# THE EXPERIMENTS WE WILL ANALYSE

## Single-cell RNA sequencing



<https://www.sciencedirect.com/science/article/pii/S1097276517300497>

<https://emea.illumina.com/science/sequencing-method-explorer/kits-and-arrays/drop-seq.html>

<https://emea.illumina.com/science/sequencing-method-explorer/kits-and-arrays/smart-seq-nanocage-cagescan.html>

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