

# **Cancer Evolution and Multistep Tumorigenesis**

***Cancer Biology I (PATH4500)***

September 24<sup>th</sup>, 2025

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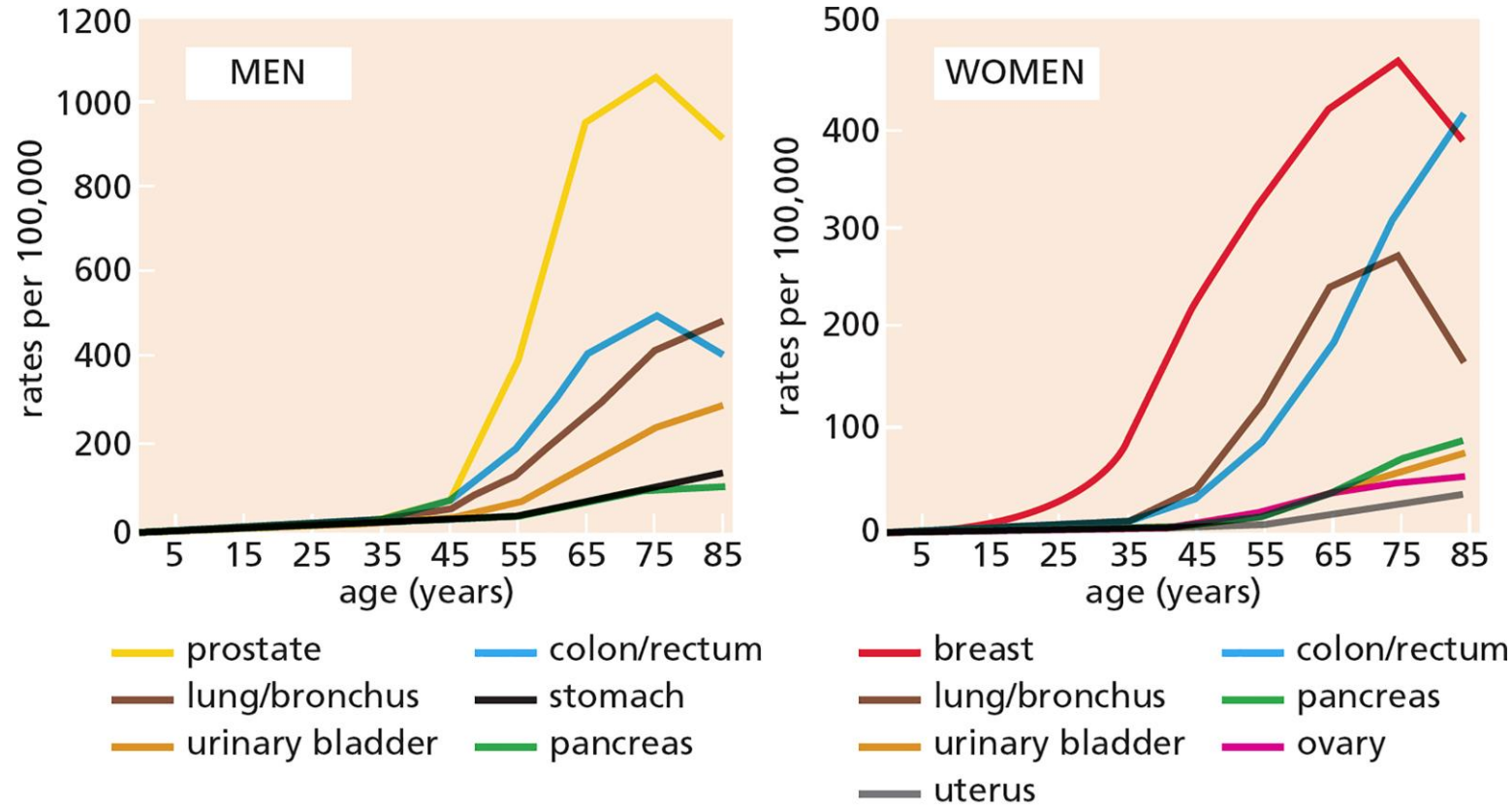
# **Cancer Evolution and Multistep Tumorigenesis**

## **-KEY LEARNING OBJECTIVES-**

- To describe the multi-step process of tumorigenesis.
- To recognize the contribution of cancer cell intrinsic mechanisms (genetic and epigenetic) and the influence of non-tumor mechanisms (e.g., tumor microenvironment, immune response).
- To examine the hallmarks of cancer.
- To identify different models of cancer evolution.

# Most human cancers develop over many decades of time

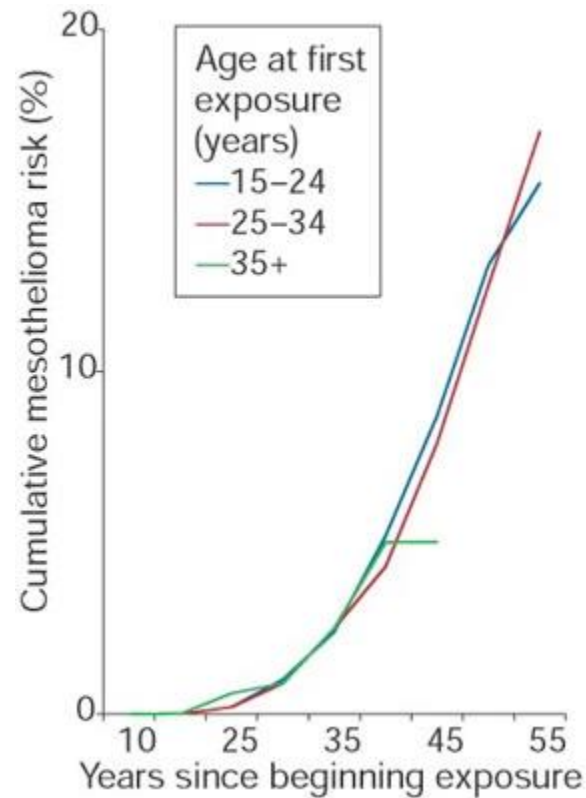
Age is a large factor in incidence of cancer



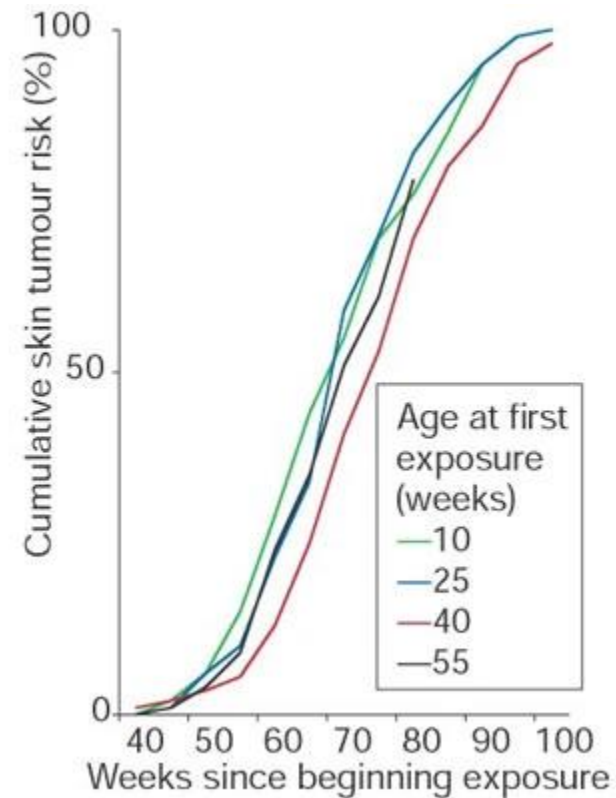
(Courtesy of W.K. Hong, compiled from *SEER Cancer Statistics Review*.)  
Copyright © 2023 W. W. Norton & Co., Inc.

Such dynamics imply a sequence of random independent events that occur at comparable frequencies over extended periods of time. The probability of these events to occur per unit of time may vary dramatically from one individual to another, being affected by inherited predisposition, diet, and lifestyle, among other variables.

# Cancer incidence and duration of carcinogen exposure



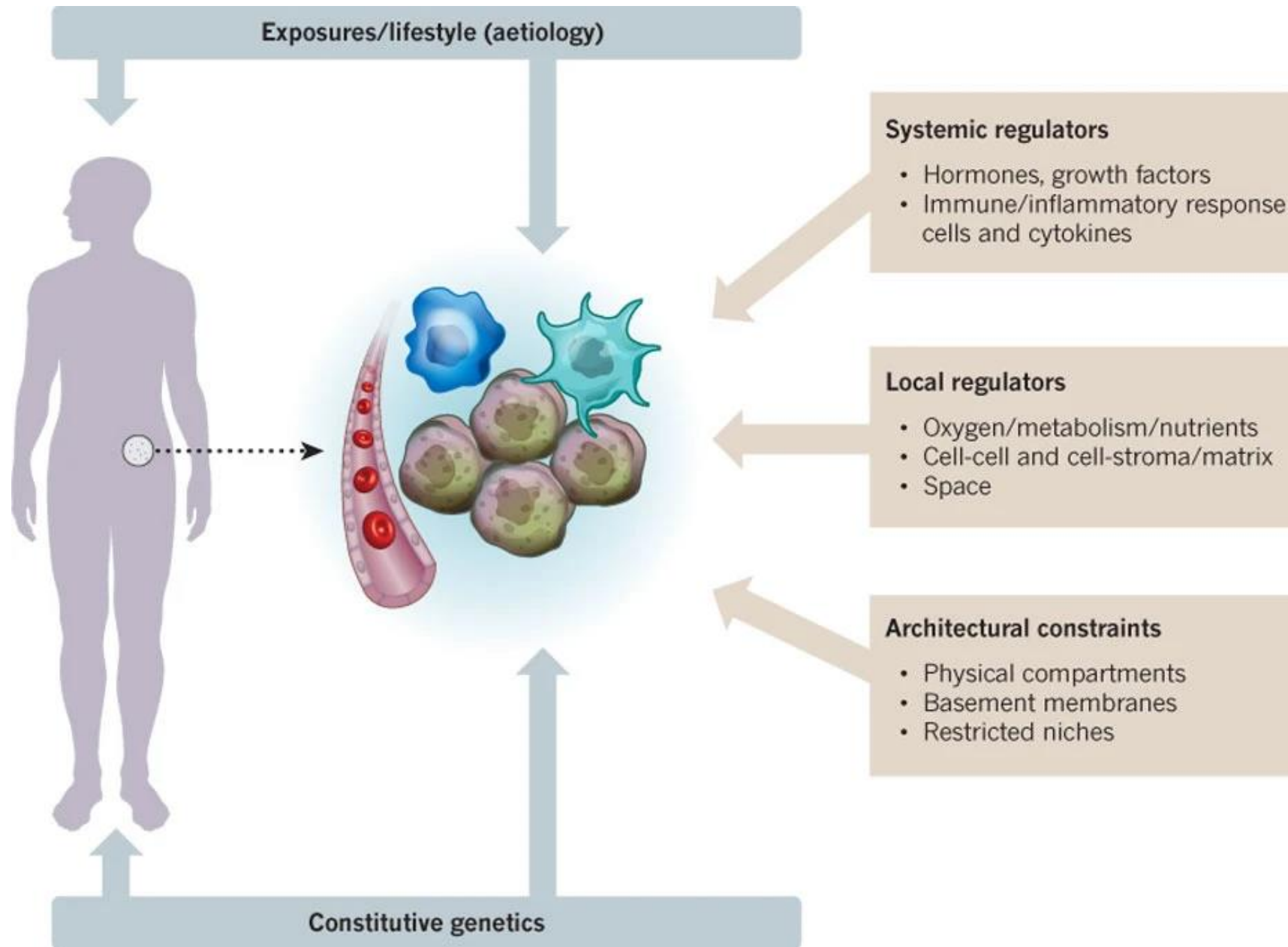
Cumulative mesothelioma risk in USA workers exposed to asbestos.



Cumulative skin tumor risk in mice treated with benzo(a)pyrene.

Malignant transformation correlates with the *duration of the exposure* rather than the age of the exposed individuals or the age when the exposure began.

# Multiple components influence cancer evolution





# Cancer development is a multi-step process -from adenoma to colon cancer-

## **Stage 0 (Carcinoma in Situ):**

This is the earliest stage of cancer. Abnormal cells are present only in the innermost lining (mucosa) and have not spread.

## **Stage I:**

The tumor has grown into the layer of tissue just below the inner lining (submucosa) or into the thick muscle layer (muscularis propria), but it has not spread to lymph nodes or distant organs.

## **Stage II:**

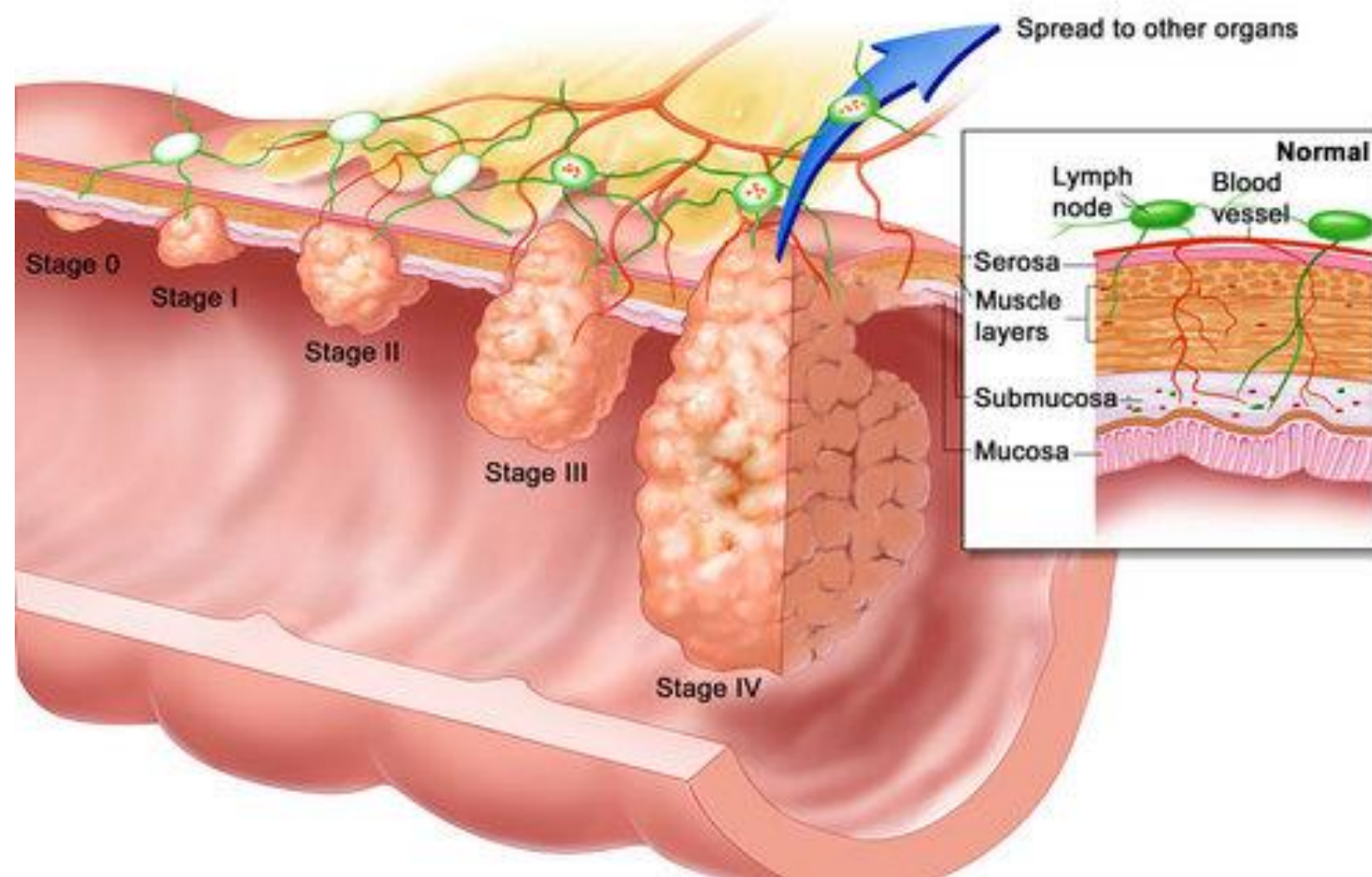
The tumor has grown through the colon wall and possibly into nearby tissues or organs, but it has not spread to lymph nodes or distant sites.

## **Stage III:**

The cancer has spread to one or more nearby lymph nodes, but there is no distant metastasis.

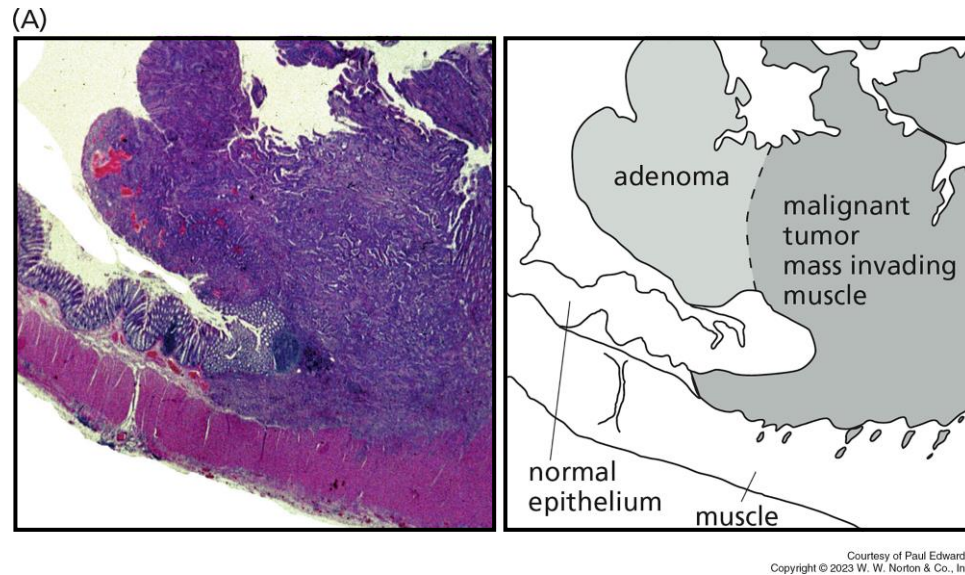
## **Stage IV:**

The cancer has spread to distant organs, such as the liver or lungs.

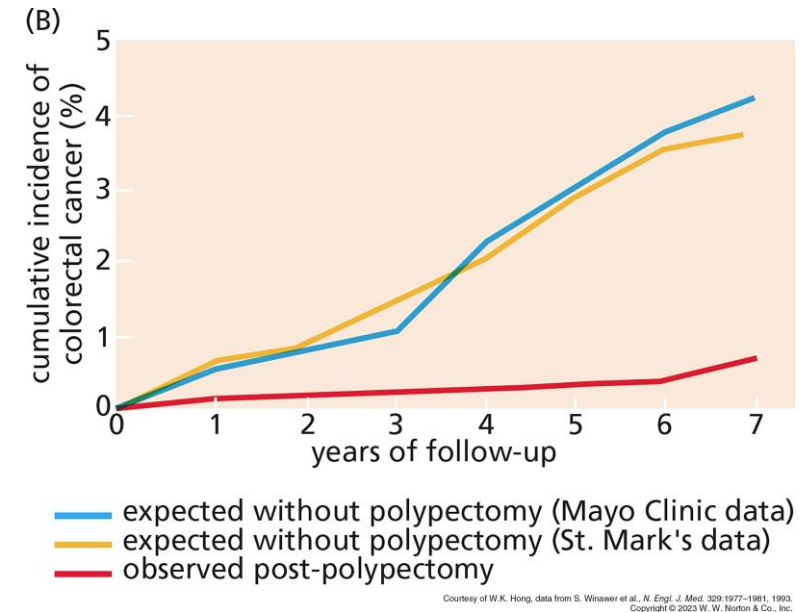


# Cancer development is a multi-step process

## -evidence for adenoma to carcinoma progression-



Occasionally, carcinomas are observed to be growing directly out of adenomas.

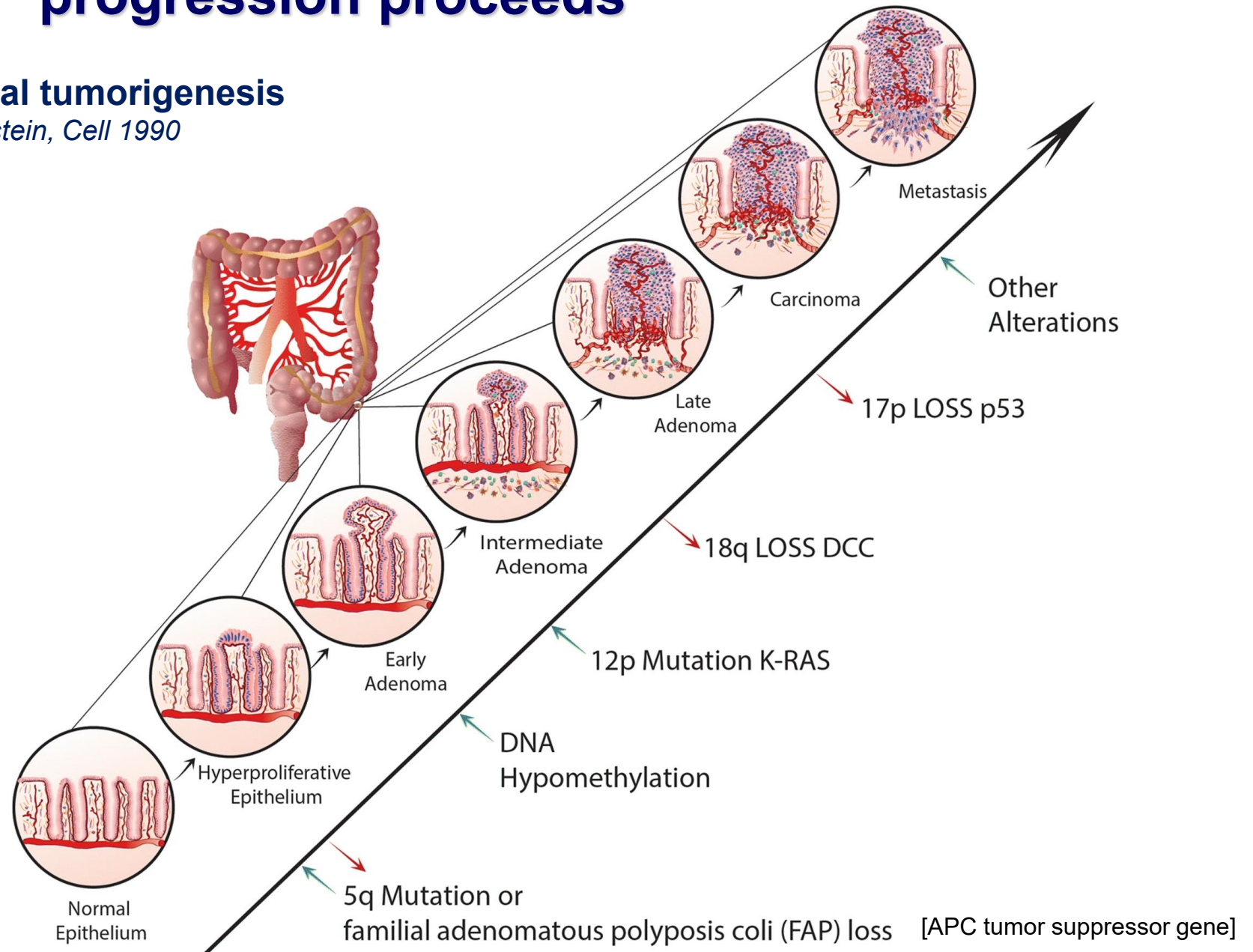


Polypectomy (surgical removal of polyps) reduces the risk of colorectal cancer.

# Cells accumulate genetic and epigenetic alterations as tumor progression proceeds

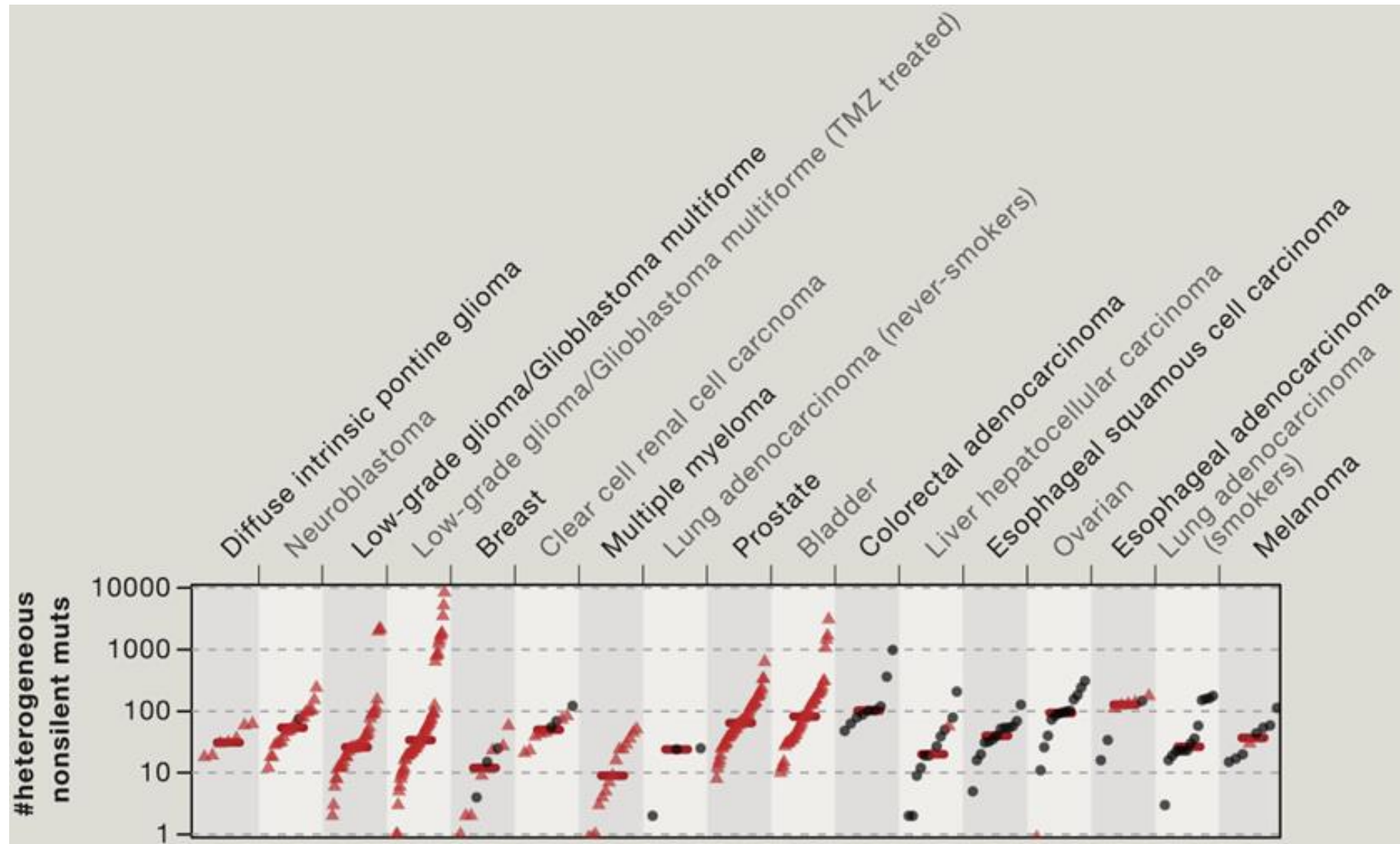
## A genetic model for colorectal tumorigenesis

adapted from *Fearon & Vogelstein, Cell 1990*

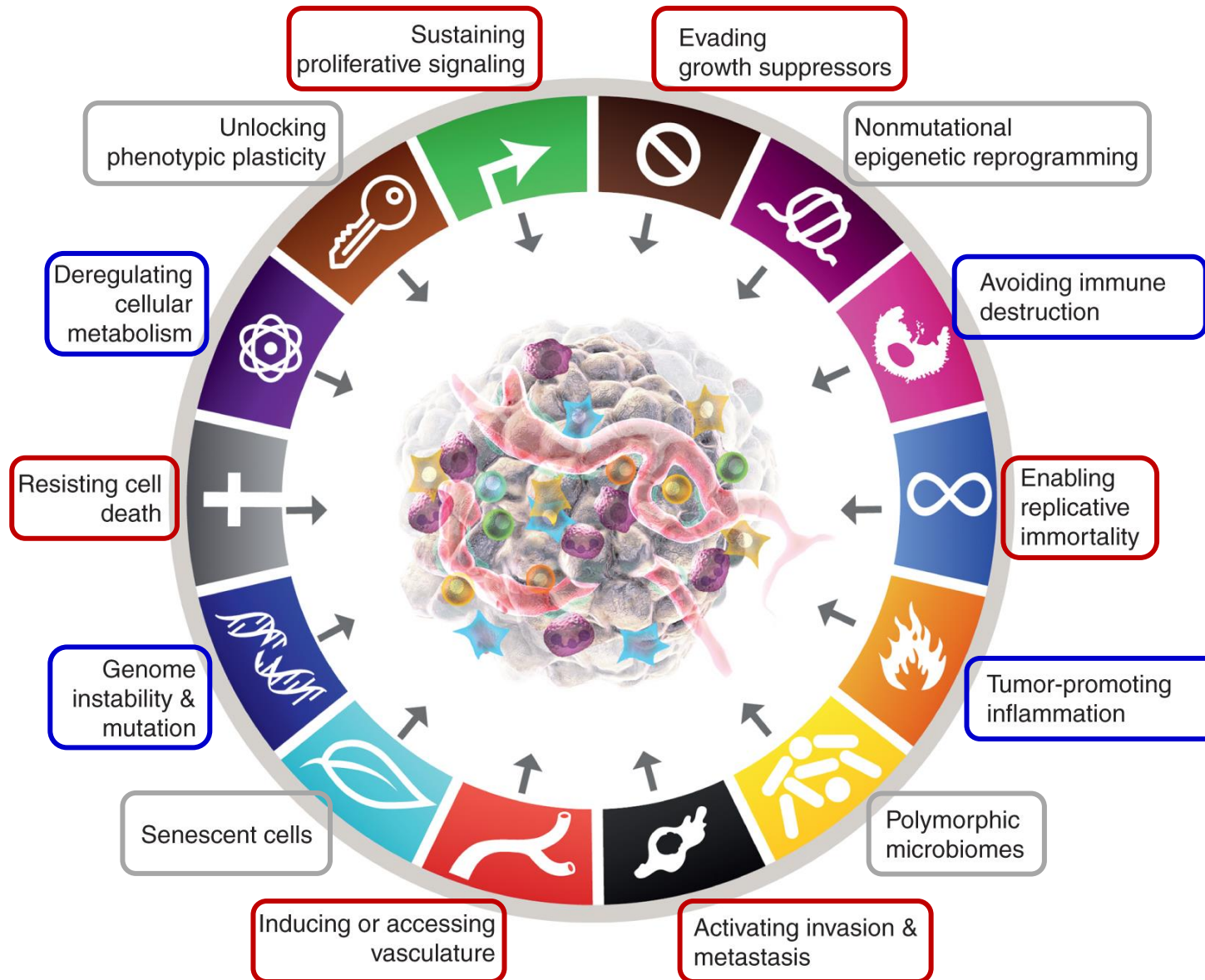




# Hundreds of non-silent mutations are detected in cancer cells



# Hallmarks of Cancer



## The Hallmarks of cancer

*Hanahan & Weinberg, Cell 2000*

## Hallmarks of cancer: next generation

*Hanahan & Weinberg, Cell 2011*

## Hallmarks of cancer: new dimensions

*Hanahan, Cancer Discovery 2022*

# **Cancer Evolution**

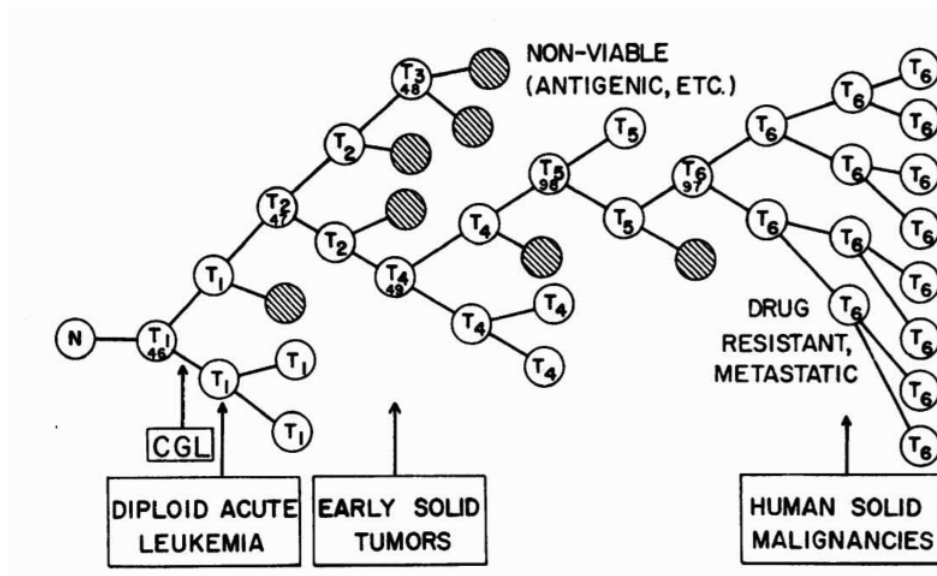
# The Clonal Evolution of Tumor Cell Populations

Acquired genetic lability permits stepwise selection of variant sublines and underlies tumor progression.

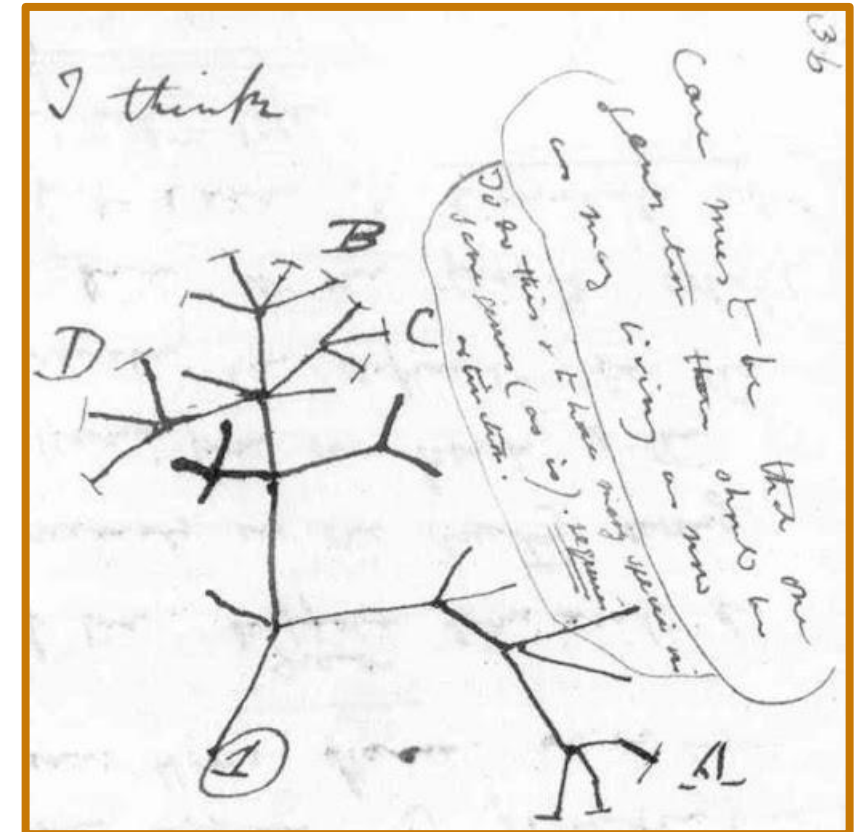
Peter C. Nowell

*Science* 1976

## Model of clonal evolution in neoplasia



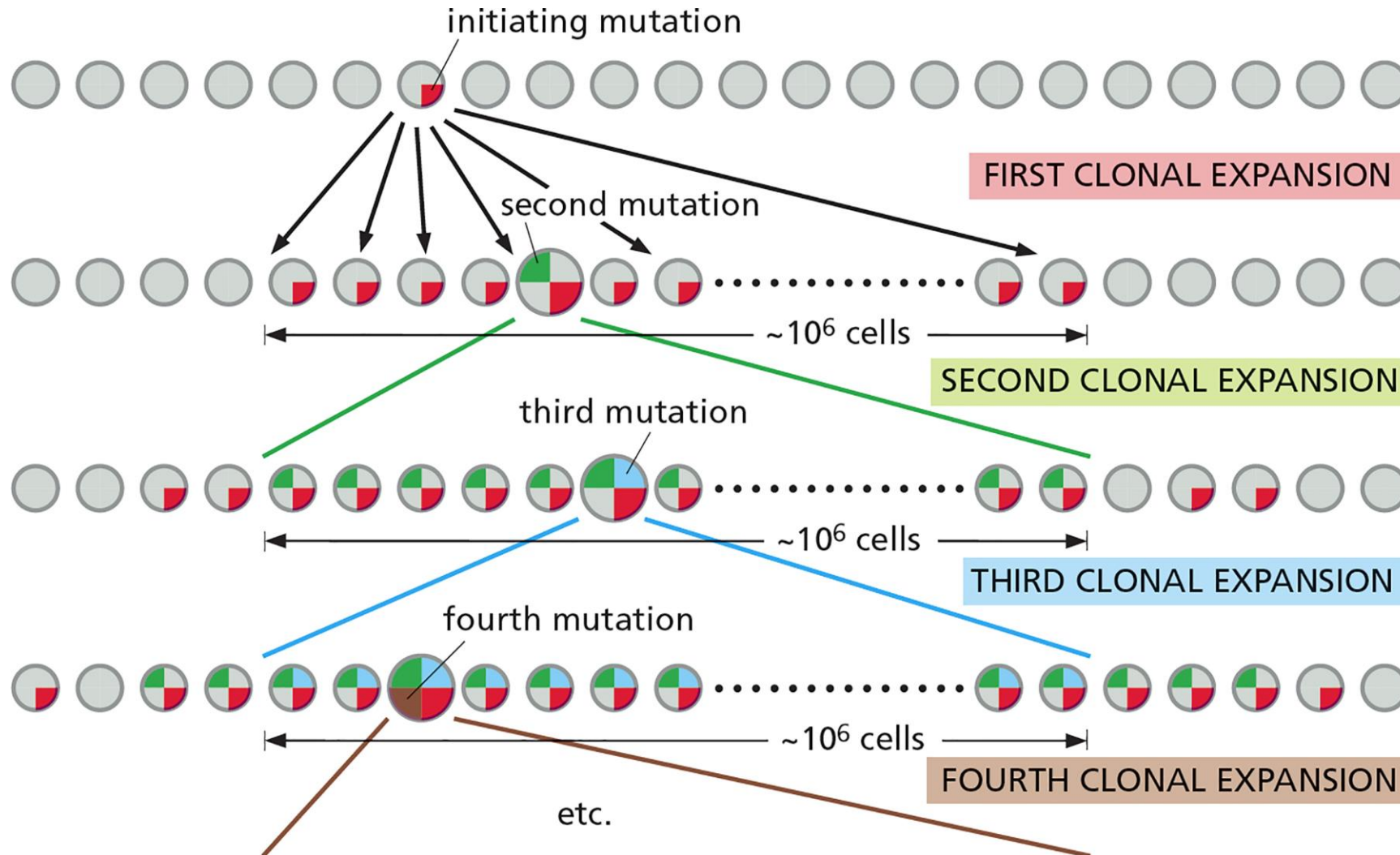
## Branching evolutionary tree of speciation (Charles Darwin, 1837)





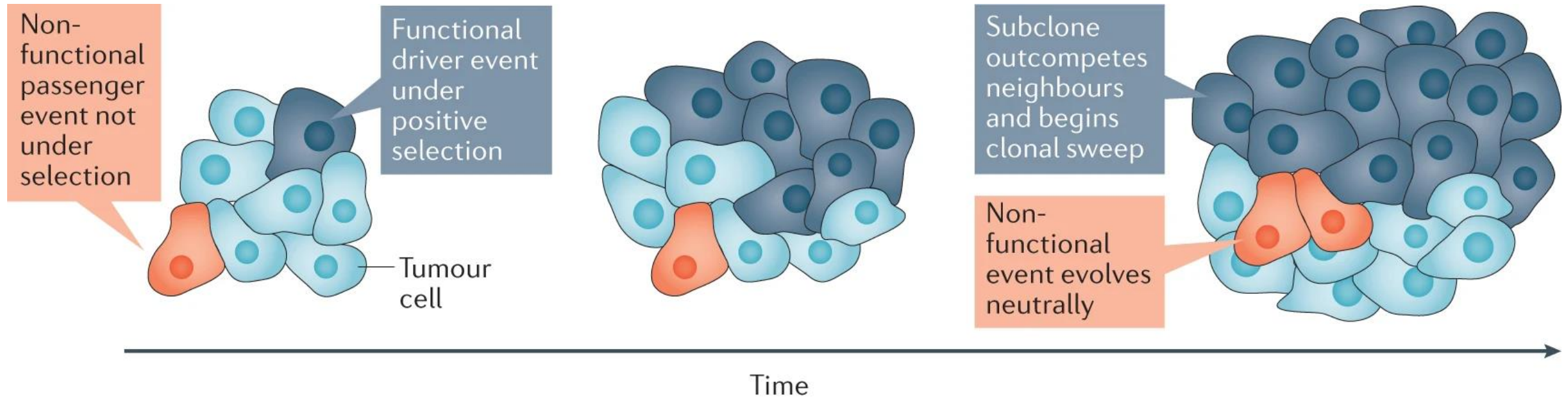
# Cancer development follows the rules of Darwinian evolution

Random mutations create genetic and phenotypic variability in a cell population, and the forces of selection may favor the outgrowth of cells with increased fitness.



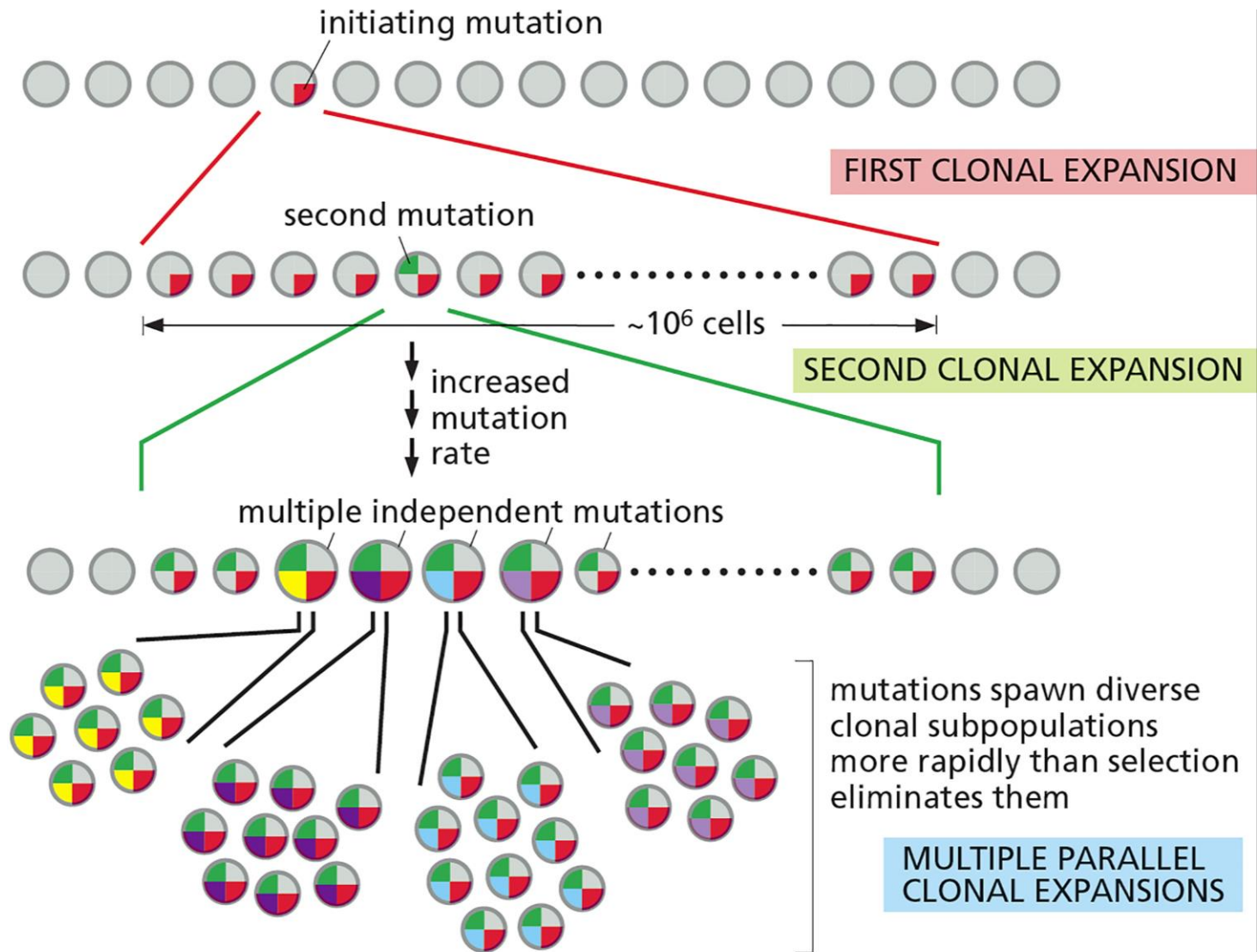


# Functional and non-functional intra-tumor heterogeneity in tumor evolution

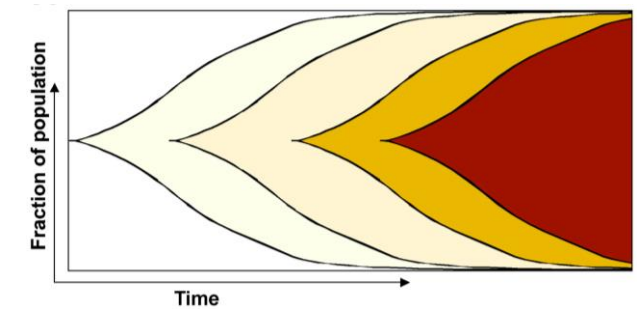


The increased rate of phenotypic variation in cancers compared with normal tissues means that new subclones arise and compete. A minority contain a driver event, such as a genetic mutation or copy number alteration, that grants a selective advantage. These subclones may grow at a faster rate than their neighbors and outcompete them in a 'selective sweep'.

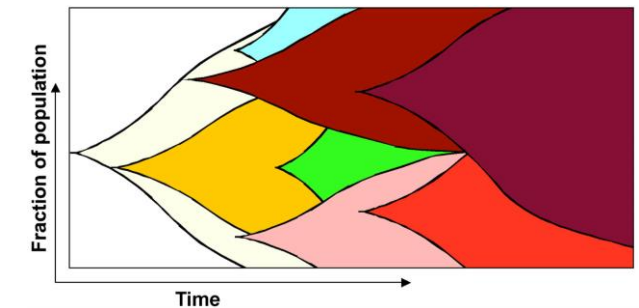
# Clonal diversification within a tumor



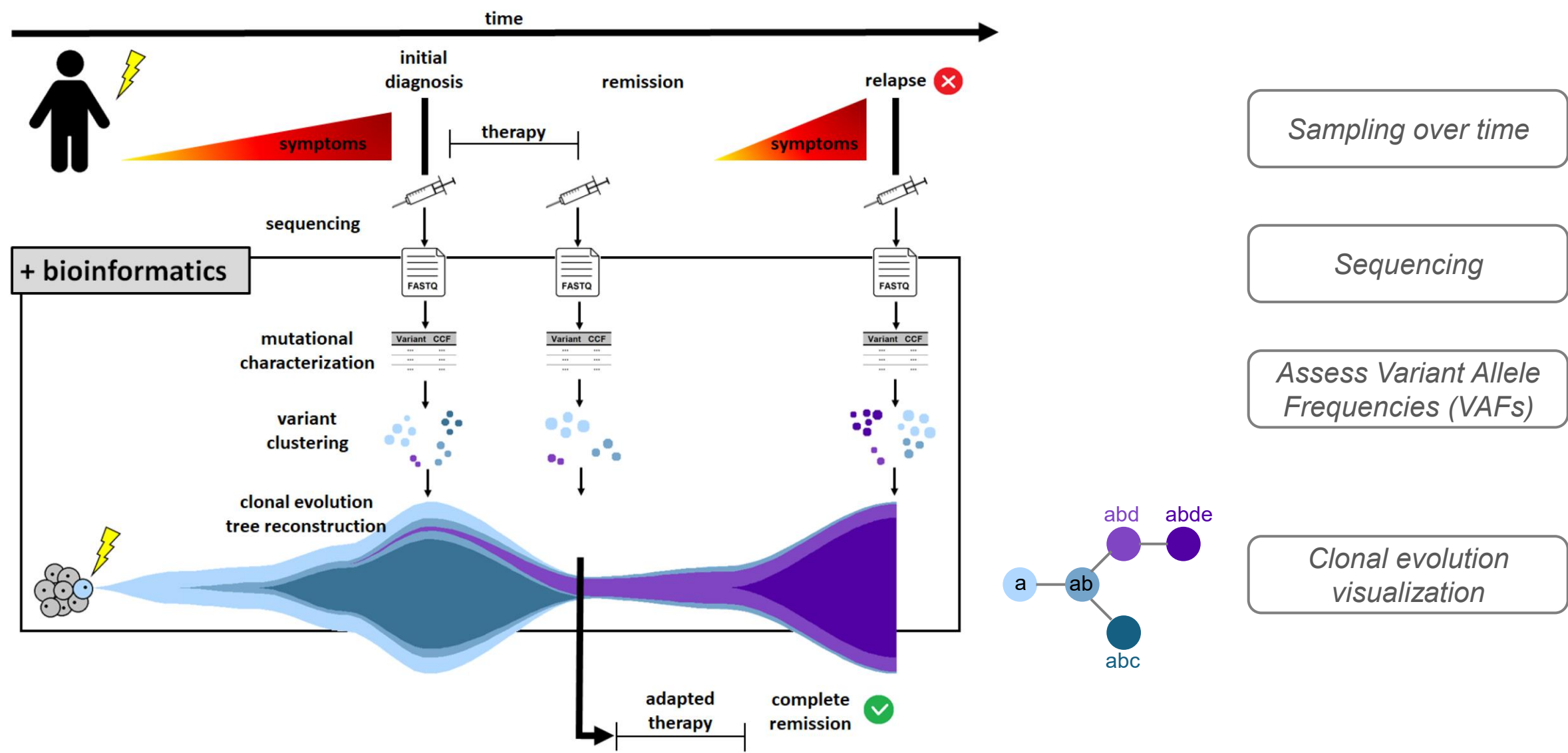
## Linear clonal succession



## Dynamic clonal diversification

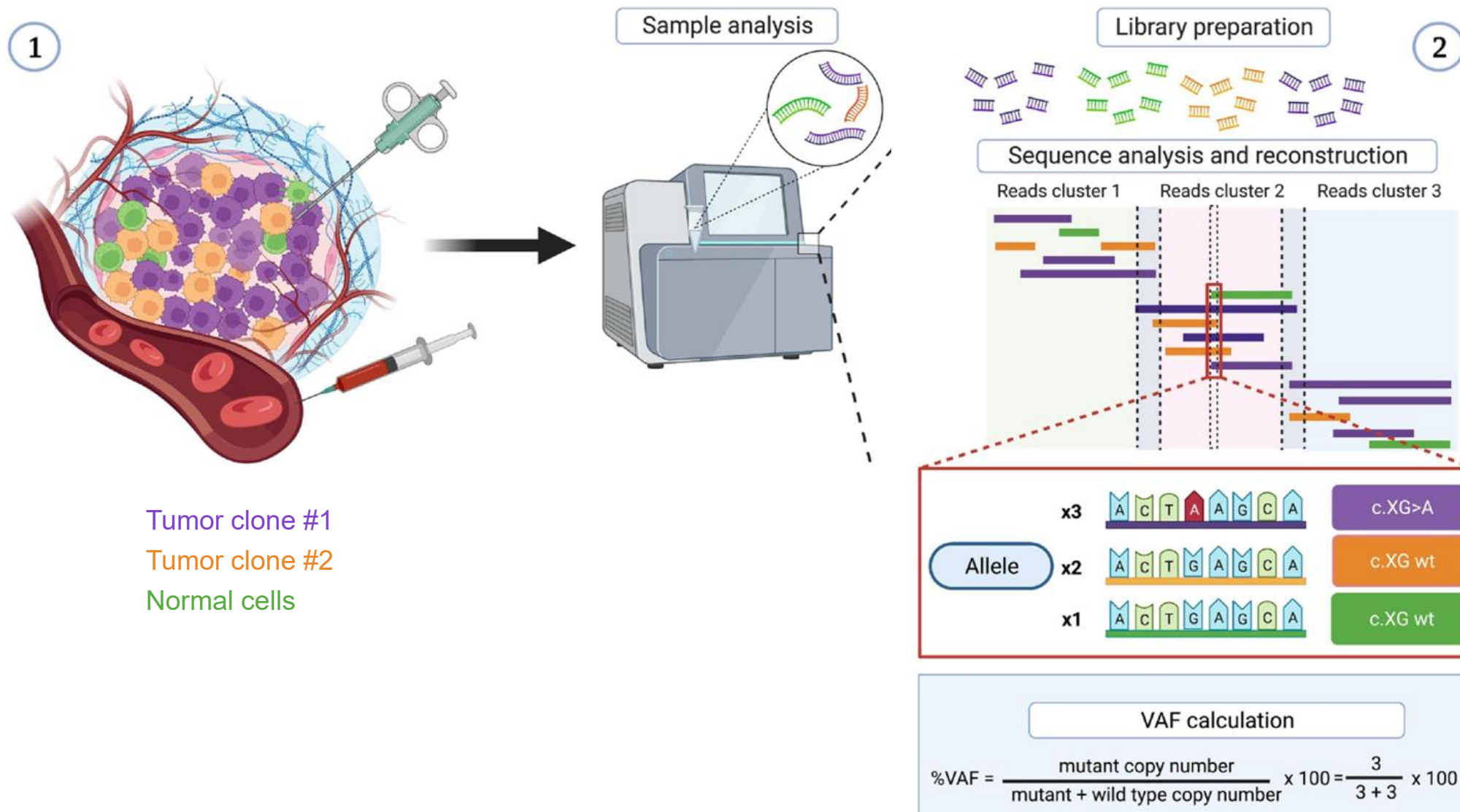


# Inferring tumor evolution and phylogenetic structure from genetic data

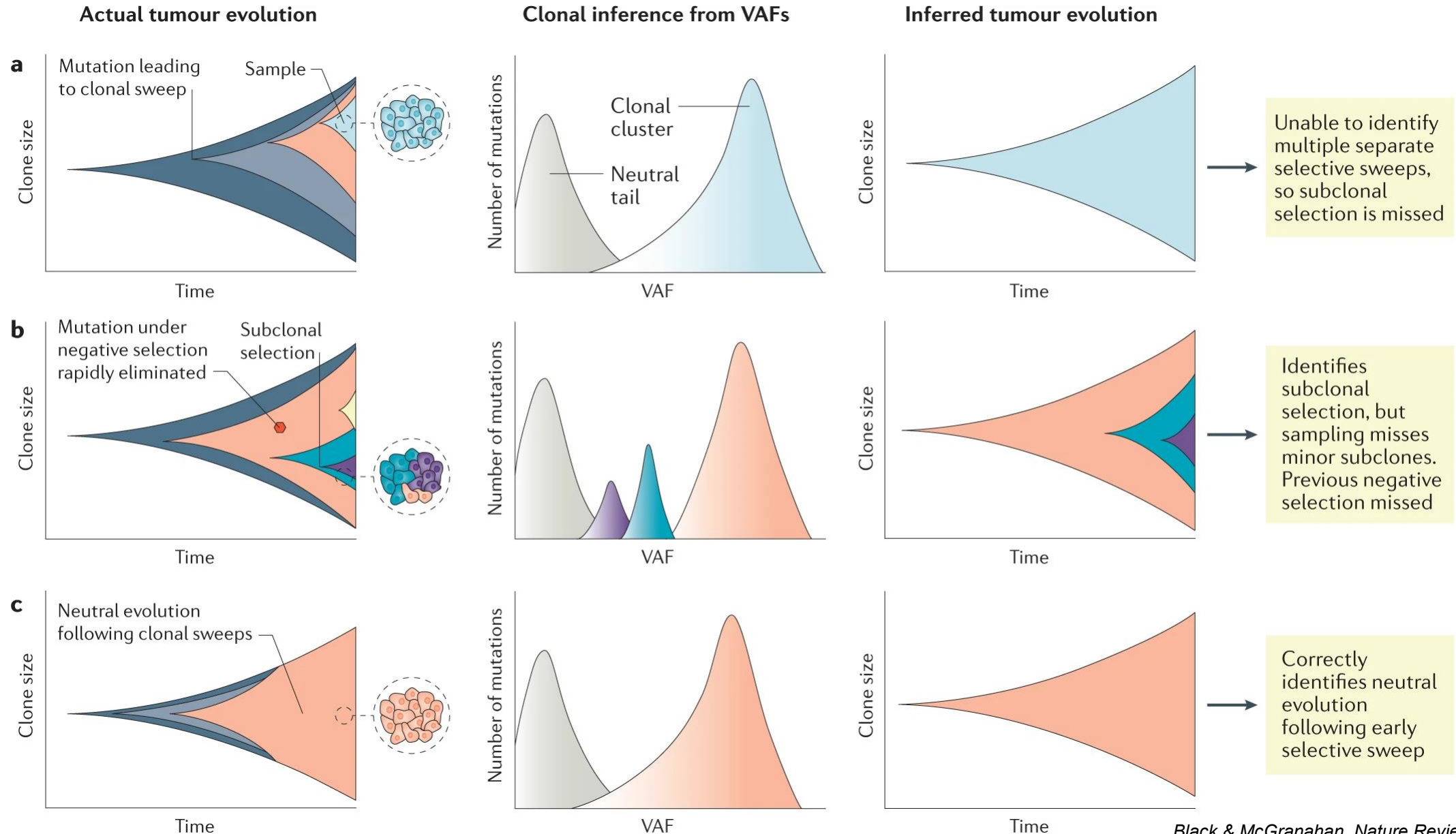




# Inferring Variant Allelic Frequencies (VAFs) of somatic mutations



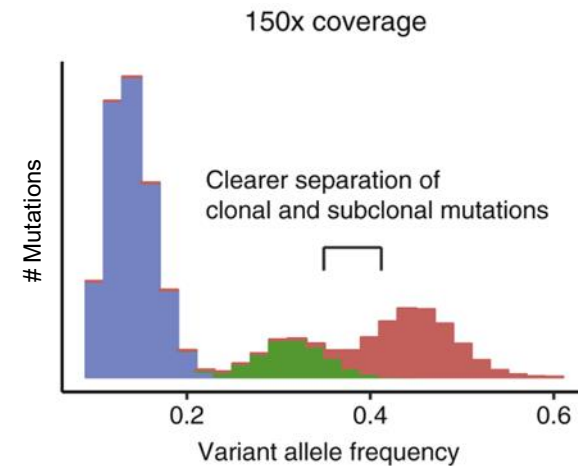
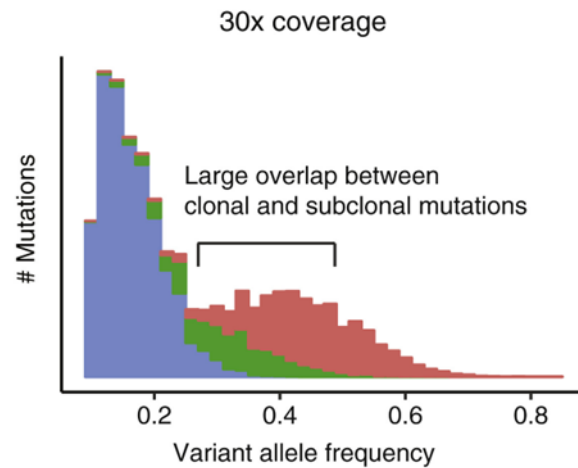
# Inferring tumor evolution from VAFs of somatic mutations



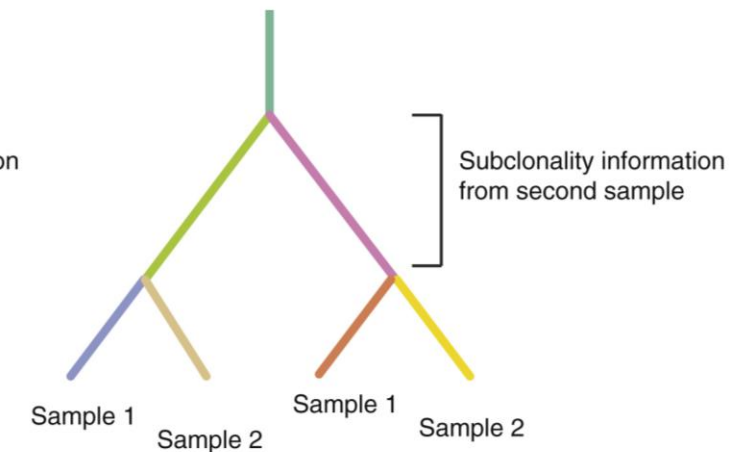
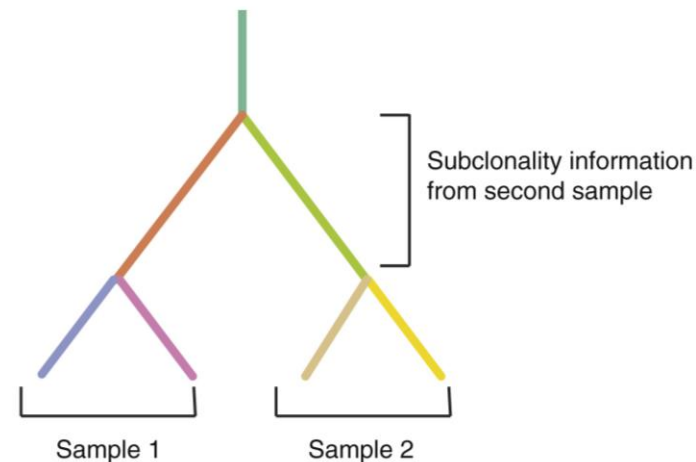
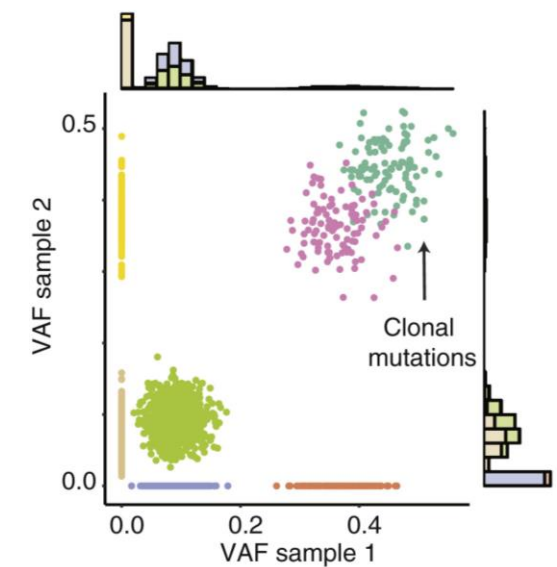
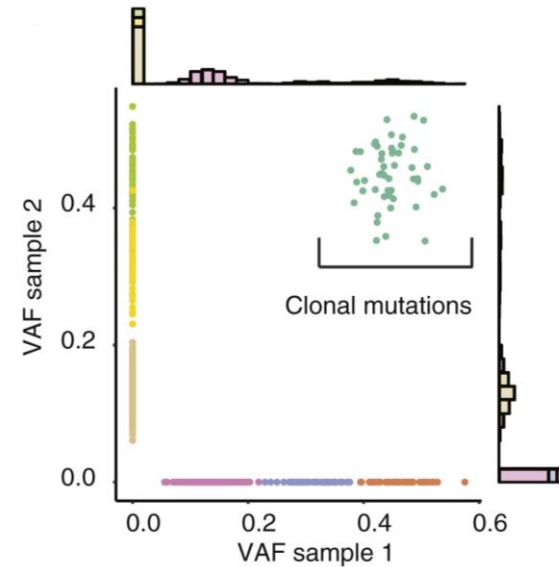


# Inferring tumor evolution and phylogenetic structure from VAFs

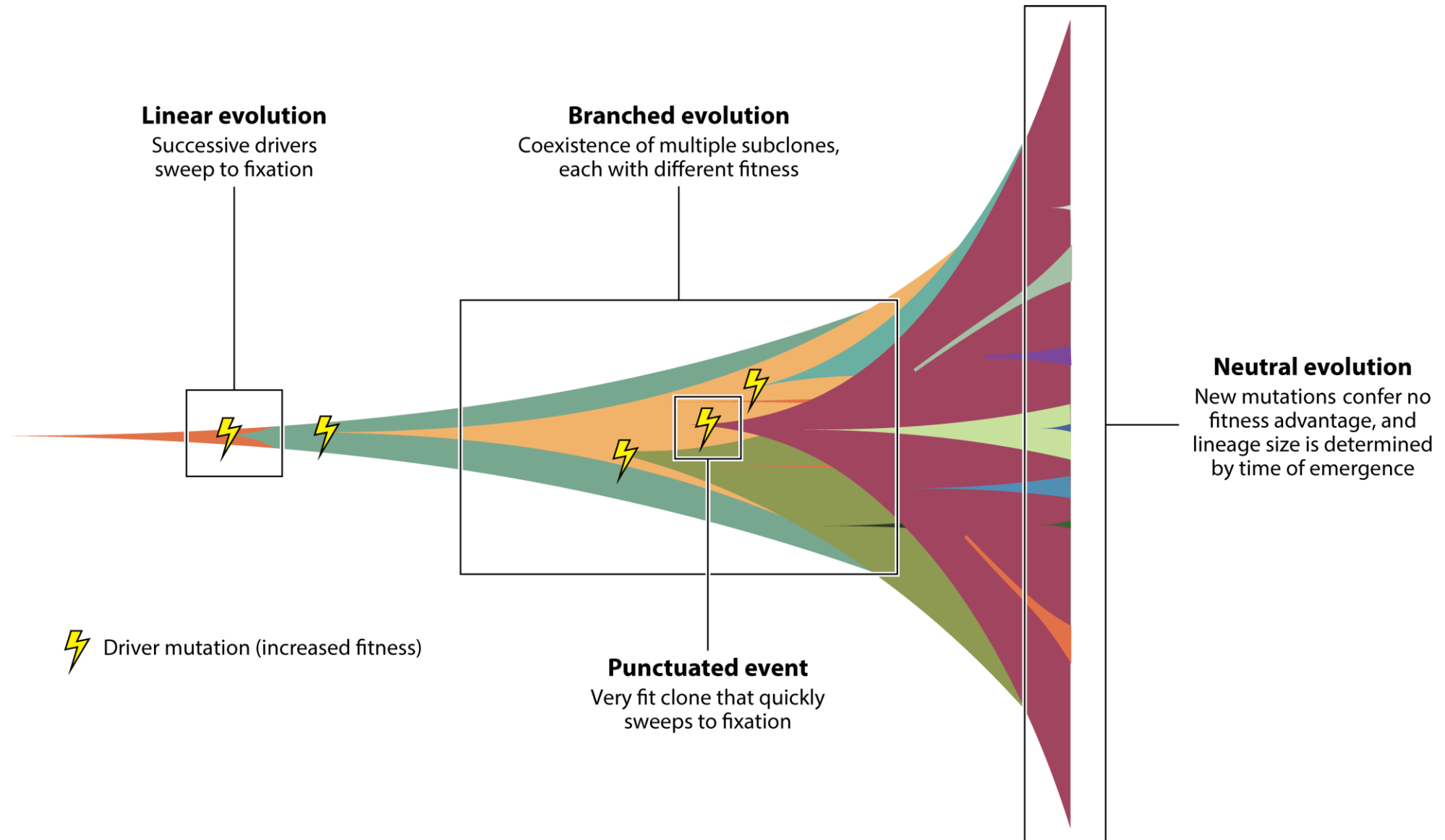
Subclonal resolution improves with higher sequencing depth



Analysis of multiple samples of the same tumor improves inference of phylogenetic structure



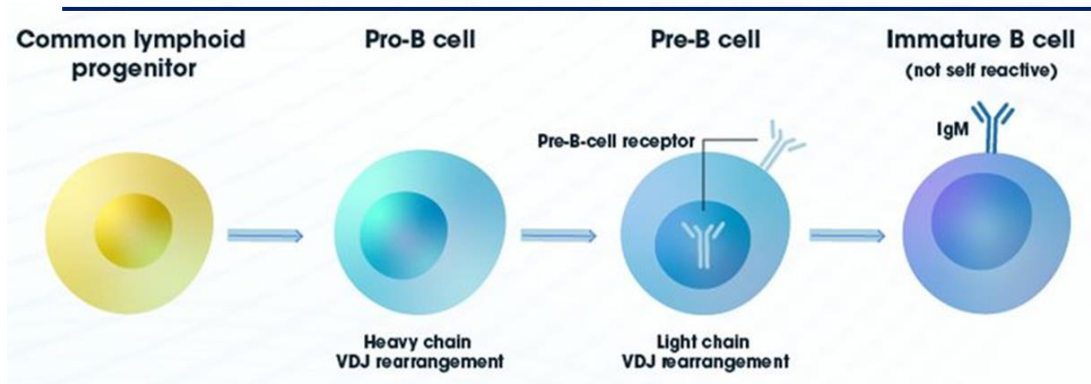
# Models of cancer evolution



# **Examples of Cancer Evolution**

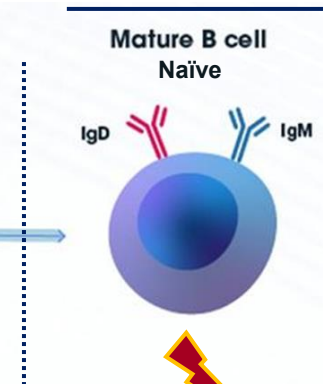
# Malignant transformation at different stages of B cell development

## BONE MARROW

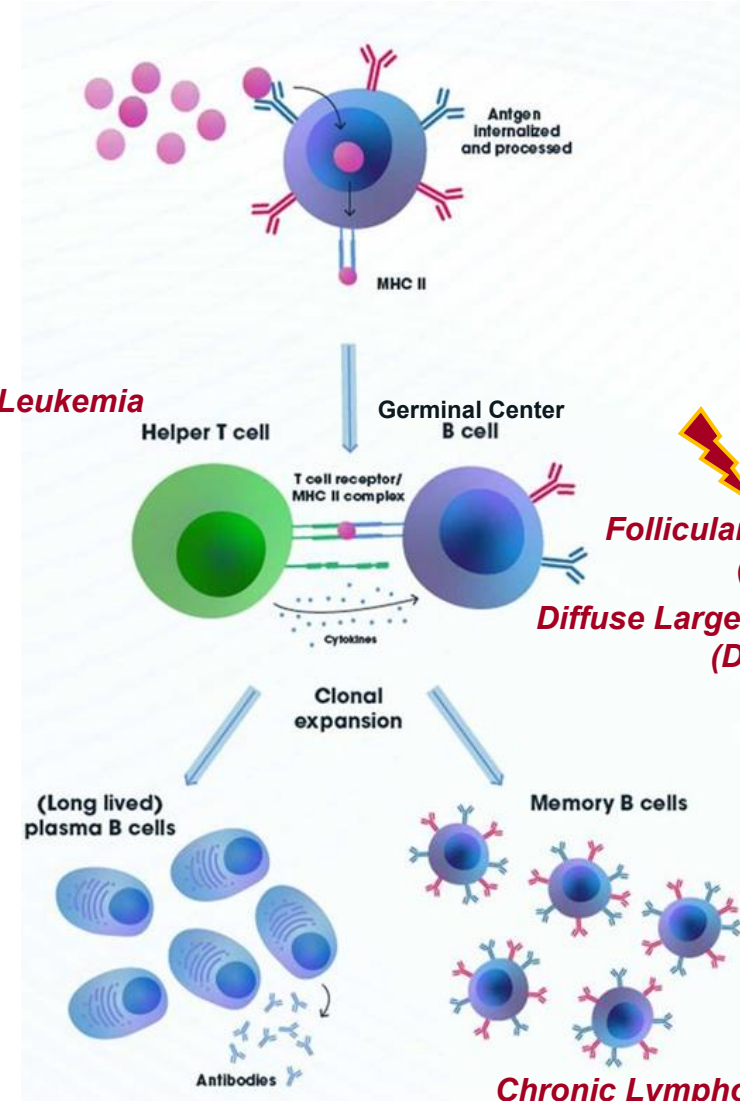


**Acute Lymphoblastic Leukemia (ALL)**

## SECONDARY LYMPHOID ORGANS



**Chronic Lymphocytic Leukemia (CLL)**



**Follicular Lymphoma (FL)**

**Diffuse Large B cell Lymphoma (DLBCL)**

**Chronic Lymphocytic Leukemia (CLL)**

# The CLL-RS transition occurs through a linear model in most cases

*Chronic Lymphocytic Leukemia  
(CLL)*

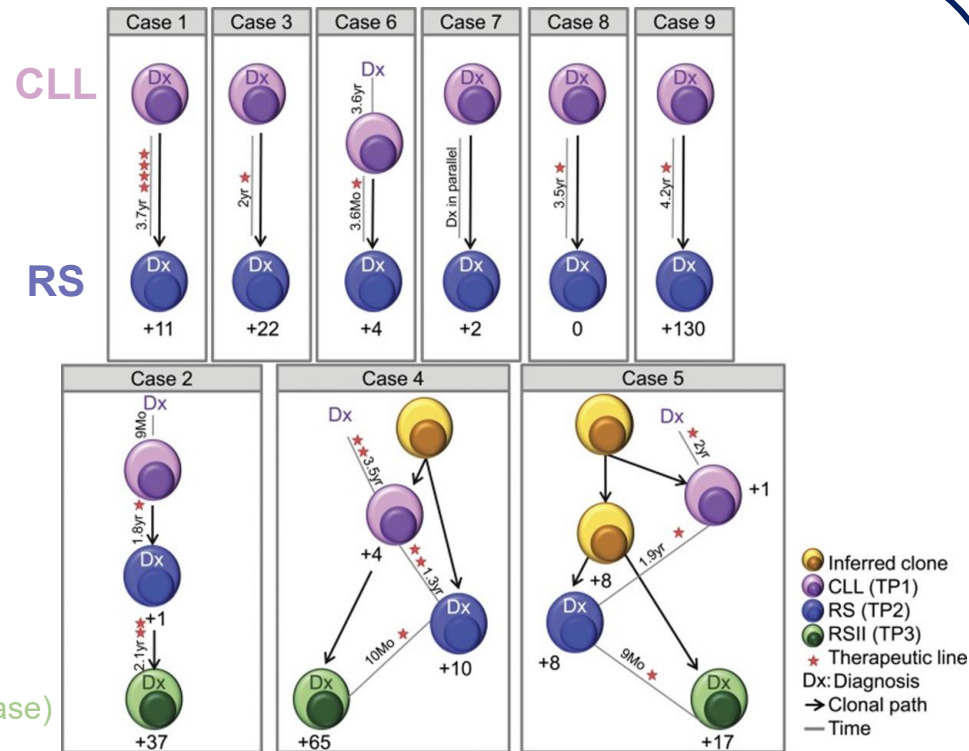
**DIAGNOSIS**  
(Time-point 1)

*Richter's Syndrome  
(RS)*

**DIAGNOSIS**  
(Time-point 2)

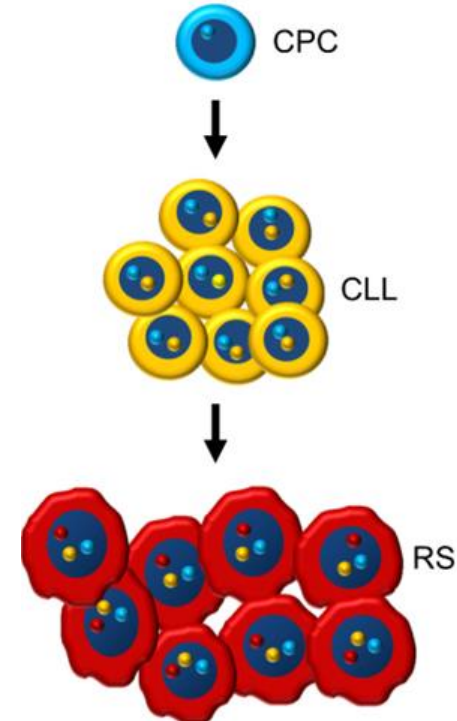
*Richter's Syndrome  
(RS leukemic phase)*

**MONITORING**  
(Time-point 3)

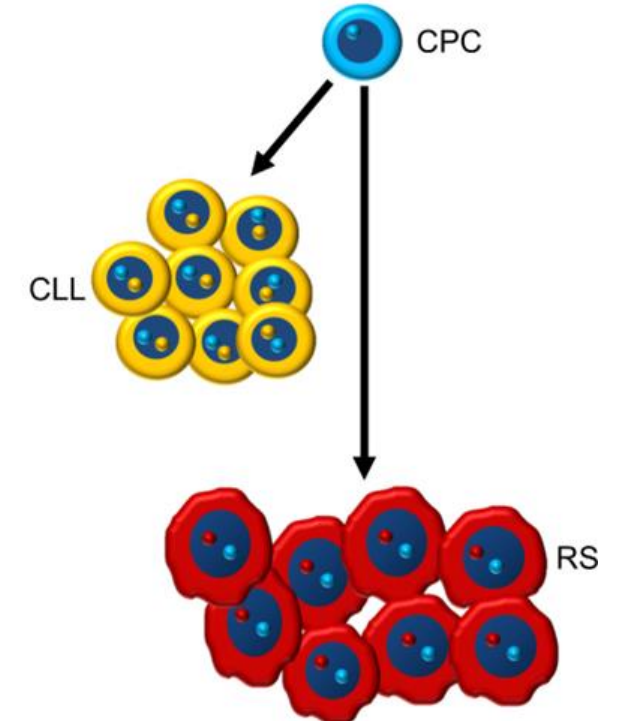


Fabbri et al., JEM, 2013

**Linear evolution  
70%**



**Branching evolution  
30%**



Rossi & Gaidano, Seminars in Oncology, 2016



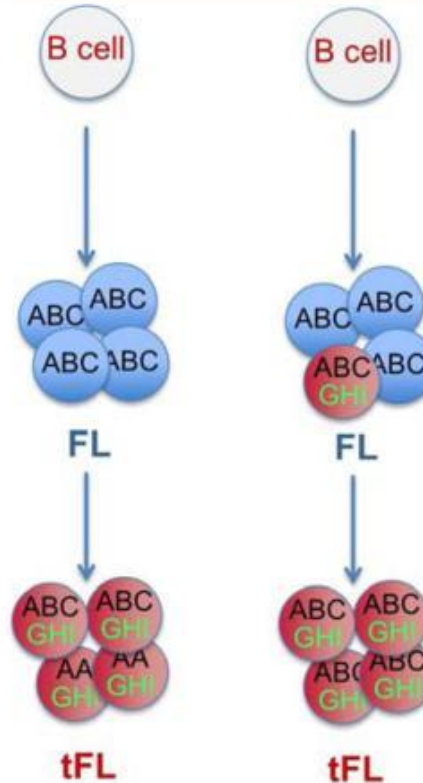
# FL transformation occurs through a branched model in most cases

**Follicular Lymphoma (FL)**

↓

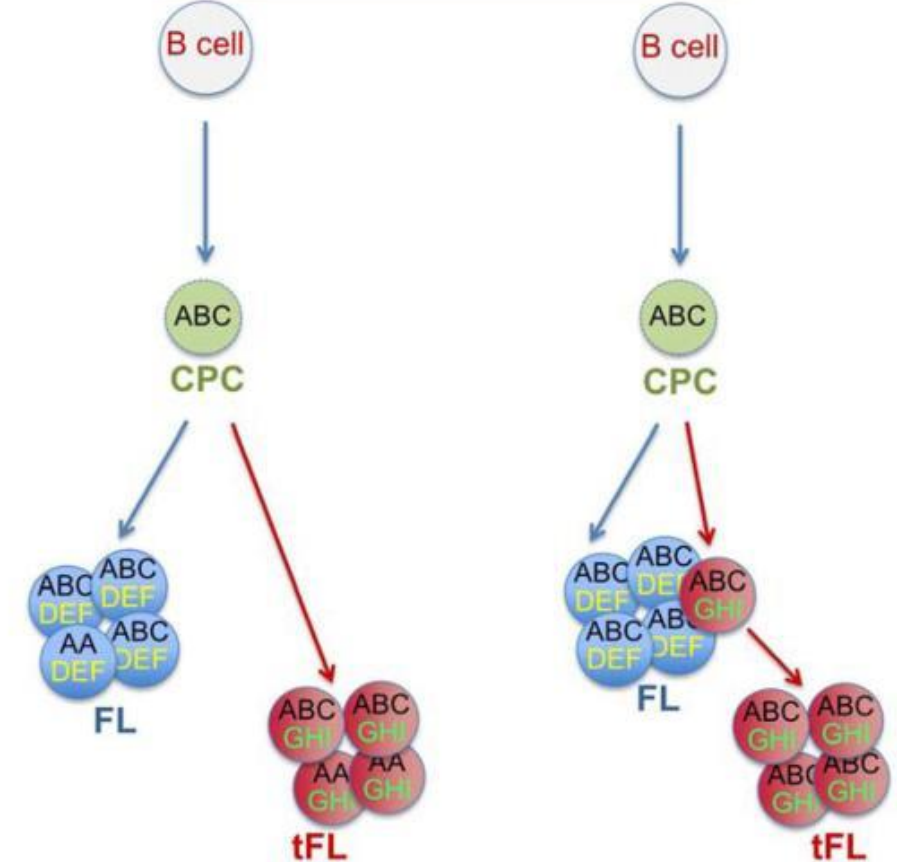
**Transformed Follicular Lymphoma (tFL)**

linear evolution  
from FL dominant clone



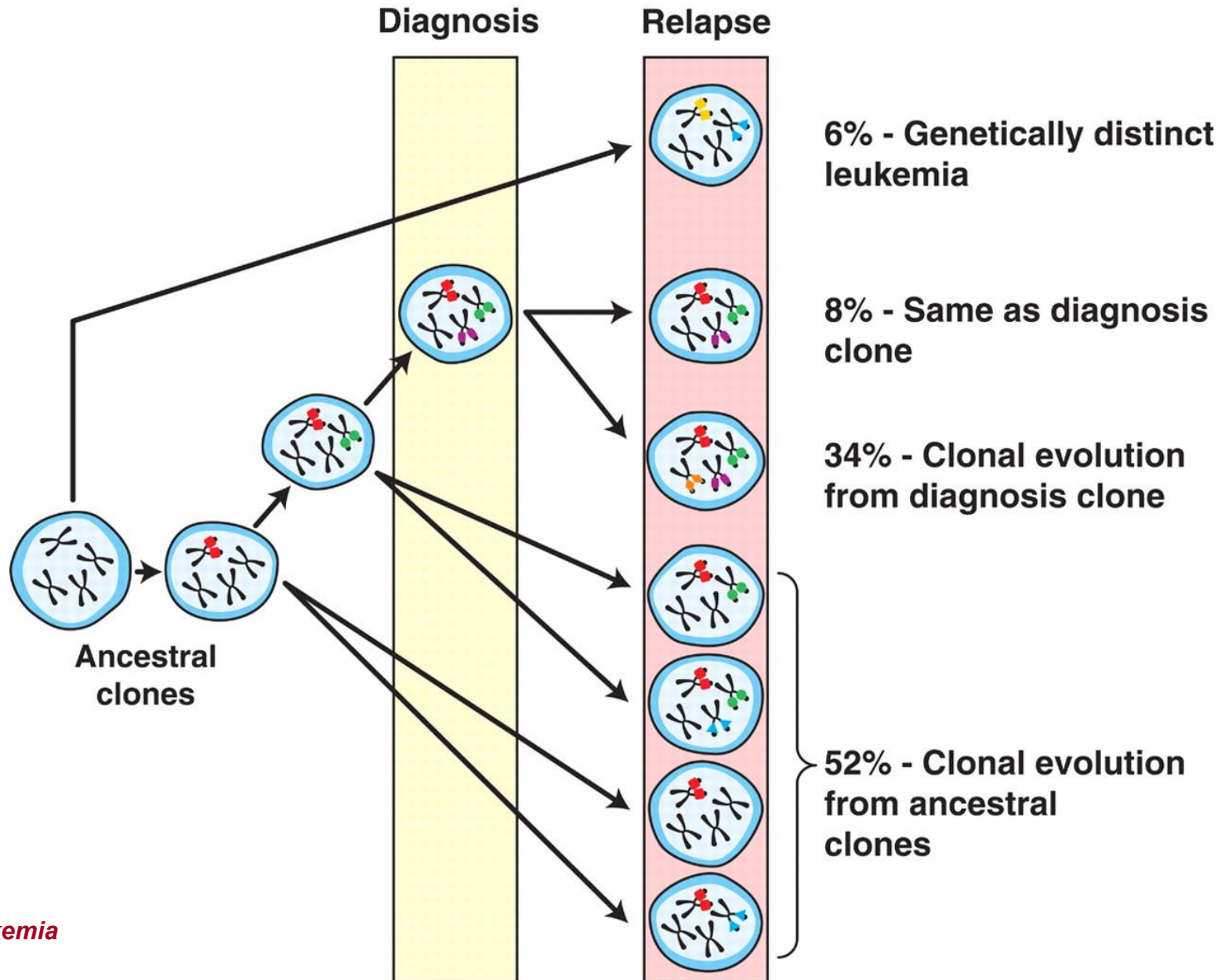
$n=2/12$  (17%)

divergent evolution  
from common progenitor cell



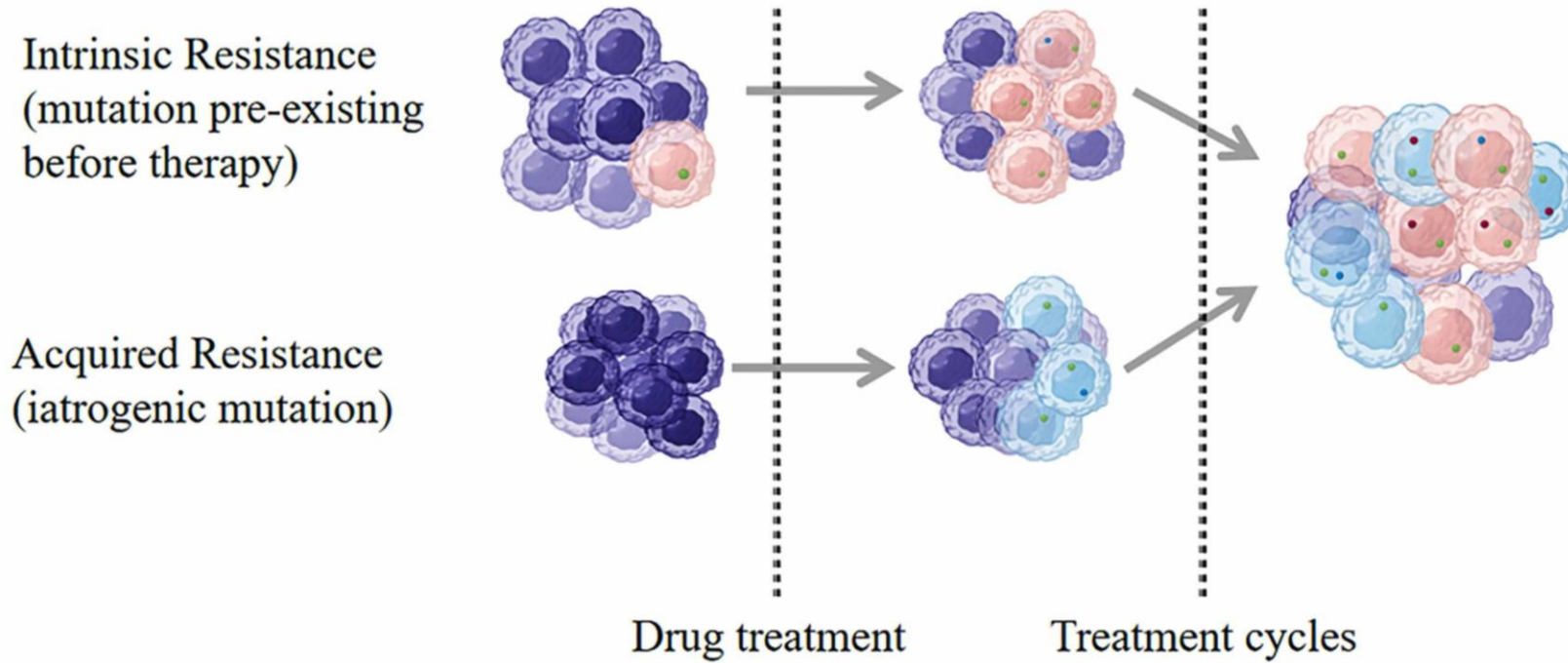
$n=10/12$  (83%)

# Clonal evolution and relapse in leukemia

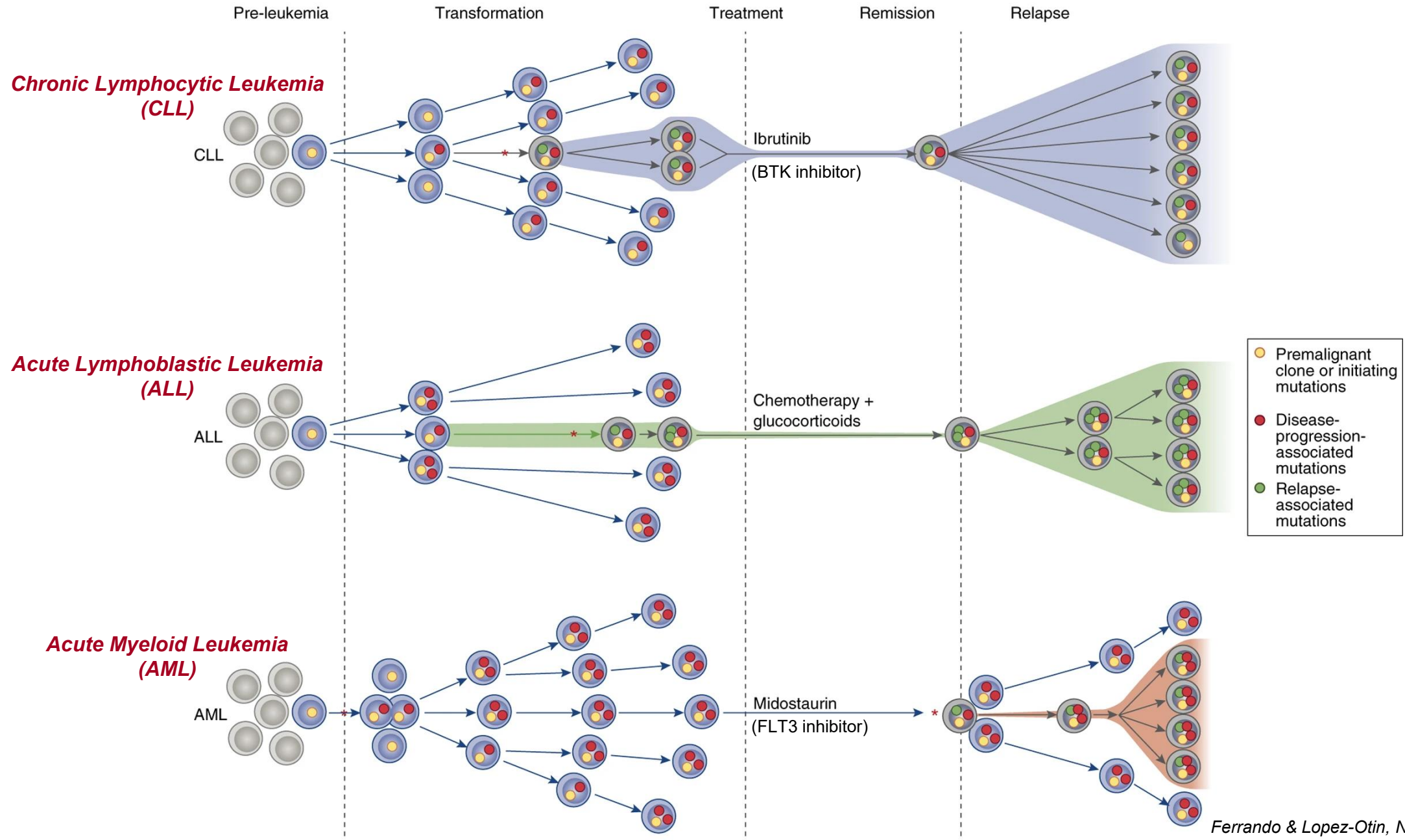


*Acute Lymphoblastic Leukemia  
(ALL)*

# Clonal diversification and development of therapy resistance



# Clonal evolution and development of therapy resistance in leukemia



# **Inferring tumor evolution from genetic data -challenges-**

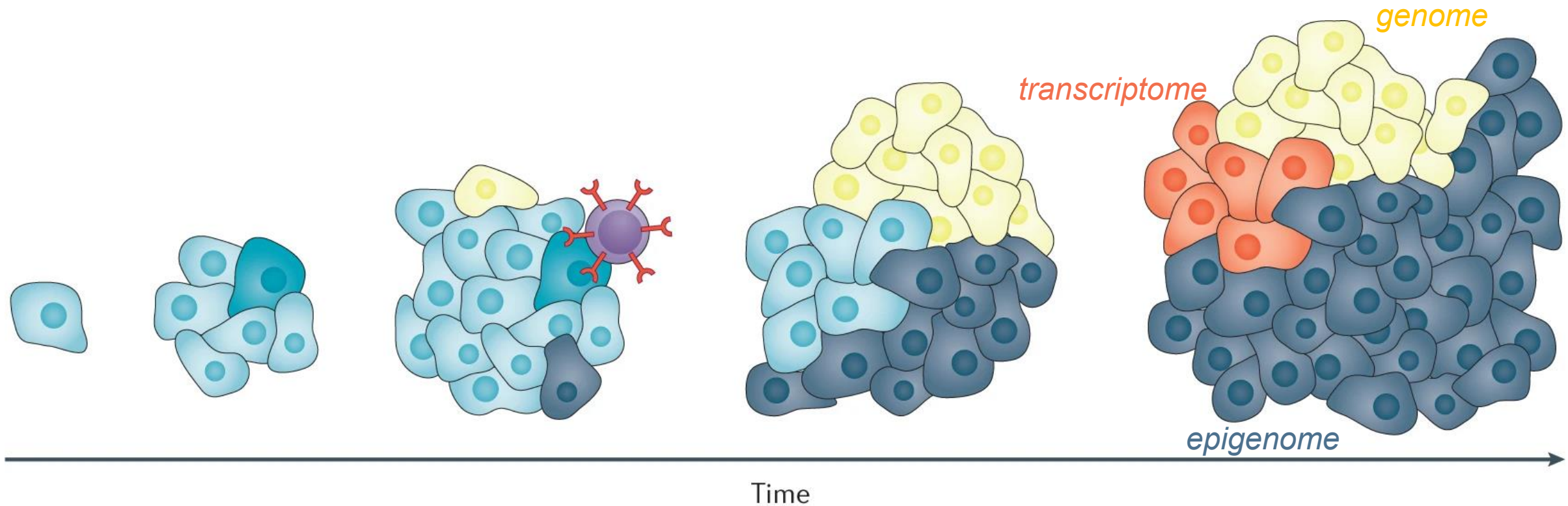




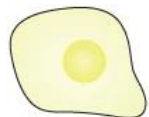
# Tumor evolution inference beyond the genomic approach

Genetic alterations are instrumental toward malignant transformation.

However, the genetic lens is only one face of the prism of forces underpinning tumor evolution.



Proliferating tumour cell



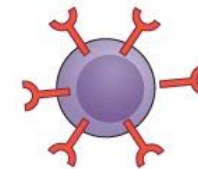
Subclonal point mutation



Subclonal epigenetic alteration



Subclonal point mutation encoding neoantigen

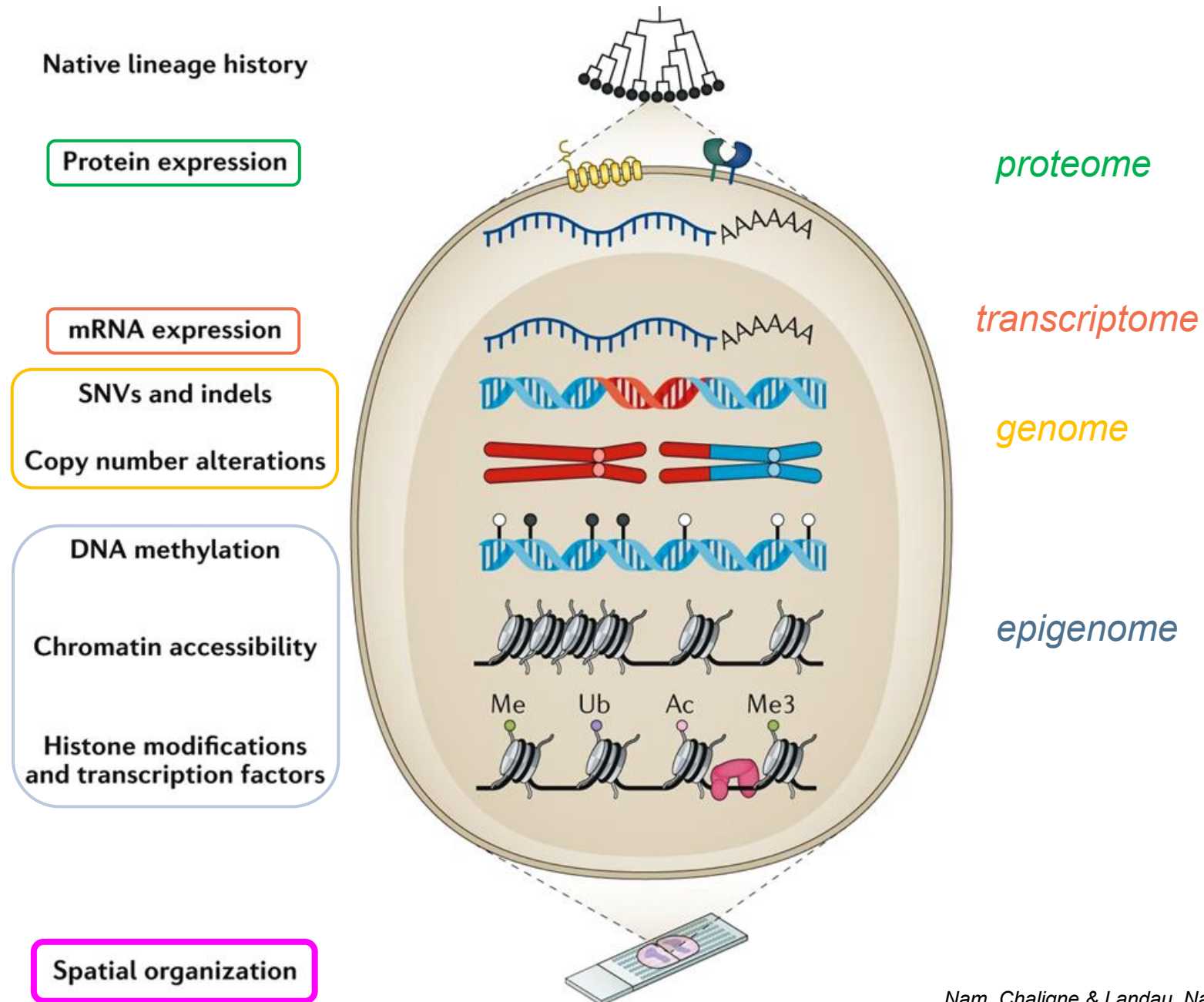


T cell with tumour antigen-specific receptors



Subclonal transcriptomic alteration

# Toward tumor evolution inference using single-cell omics data



# Cancer Evolution and Multistep Tumorigenesis

## -CONCLUSIONS-

- ❖ Tumorigenesis: a multi-step process relying on the acquisition of alterations (genetic and epigenetic) which affect multiple distinct regulatory circuits within cells and systems and function in a complementary fashion to create the neoplastic phenotype.
- ❖ The neoplastic phenotype can be achieved through multiple means and distinct alterations, however several hallmarks are shared across tumors, including: 1) ability to proliferate indefinitely; 2) reduced susceptibility to cell death; 3) acquisition of invasiveness and metastatic ability; 4) promoting angiogenesis; 5) ability to evade the immune system.
- ❖ Multi-step tumor progression can be depicted as a form of Darwinian evolution occurring within tissues. However, the contribution of epigenetic mechanisms and the rapid acquisition of genetic changes require to modify the linear Darwinian evolution model to incorporate complex branched evolution.

# Suggested Readings

## Clonal evolution in cancer.

Greaves M, **Maley CC**. Nature. 2012 Jan 18;481(7381):306-13. doi: 10.1038/nature10762.PMID: 22258609

## Genetic and non-genetic clonal diversity in cancer evolution.

Black JRM, **McGranahan N**. Nat Rev Cancer. 2021 Jun;21(6):379-392. doi: 10.1038/s41568-021-00336-2. Epub 2021 Mar 16.PMID: 33727690