

# “Cancer stem cells, evolution and heterogeneity within tumours”

**Brian Huntly**

*Stem Cell Institute and Dept of Haematology, University of Cambridge  
Co-Lead Haem Malignancies Programme  
and Honorary Consultant Haematologist, CUH*

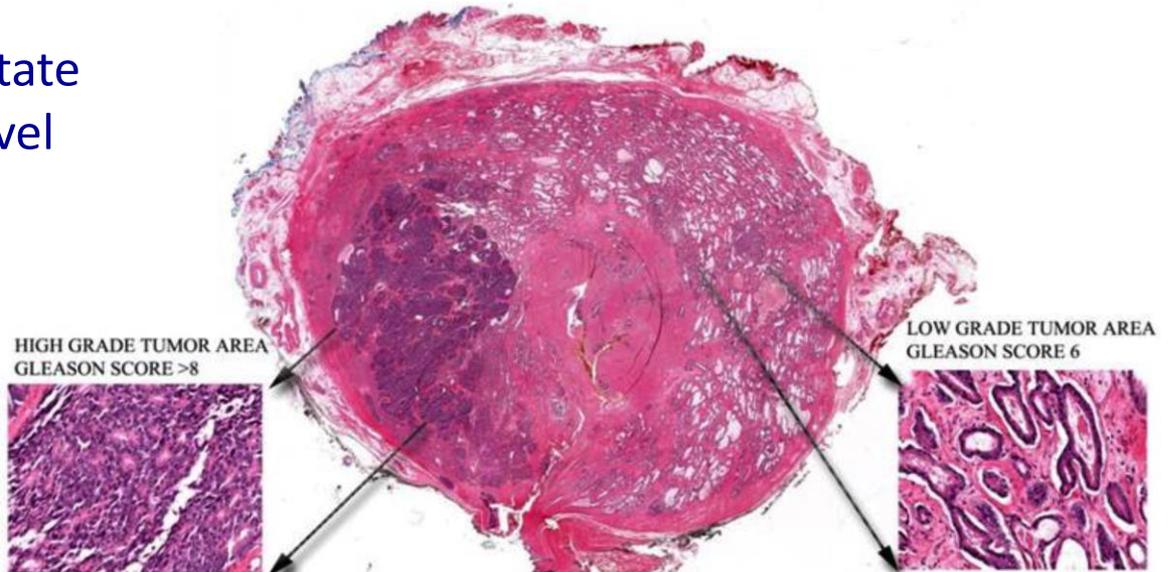
**Lecture series in Cancer Biology  
Thursday 28<sup>th</sup> November 2019**

- **Context and definitions**
- Tumour hierarchy: the cancer stem cell hypothesis and its implications for tumour biology (VERTICAL/DEVELOPMENTAL CONTEXT)
- Pre-malignancy – overt and covert premalignant states and early tumour evolution (HORIZONTAL/TIME CONTEXT)
- Tumour evolution under selective pressures:
  - Relapse
  - Metastases
- Clinical implications: Cancer stem cells  
Targeting tumour heterogeneity

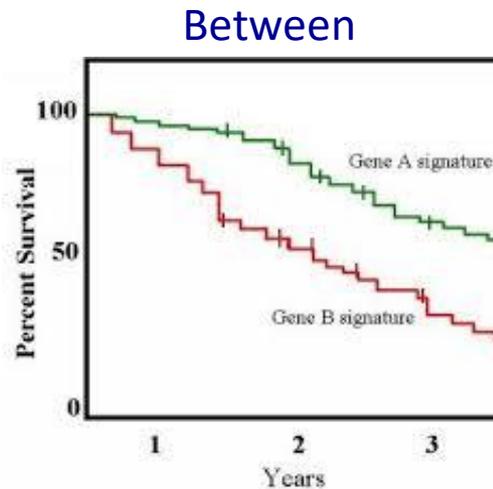
# Levels of heterogeneity

## Heterogeneity of phenotype- Between tumours and within tumours

- Visual – Malignant prostate  
Macroscopic level



- Behaviour – response to drugs

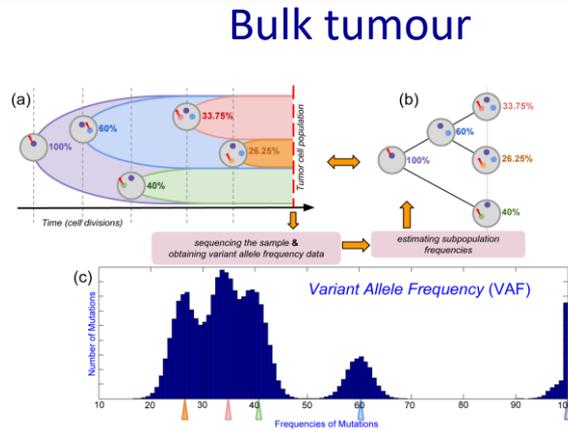


Within

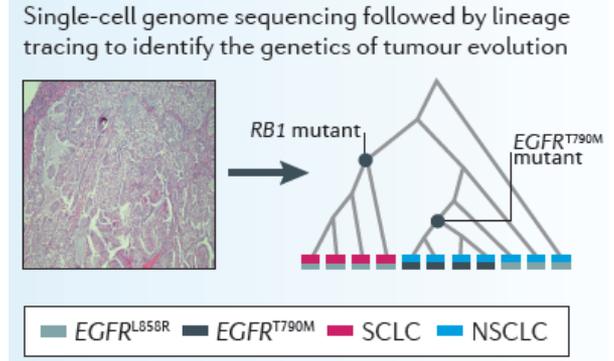
Chemo-resistant  
relapse

# Levels of heterogeneity

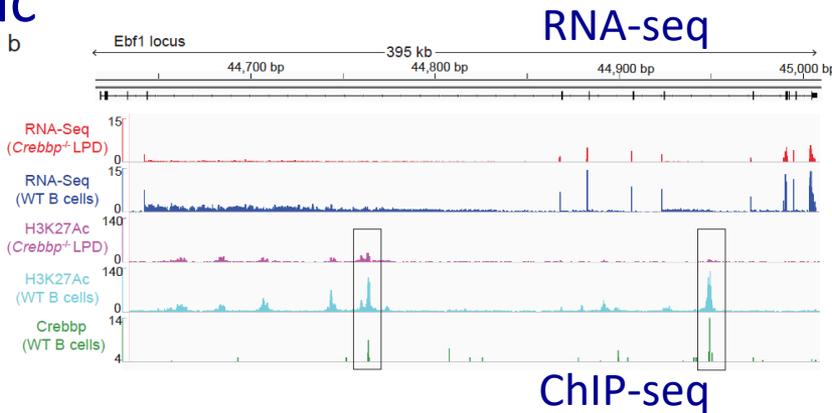
- Genetic



## Single cell analysis



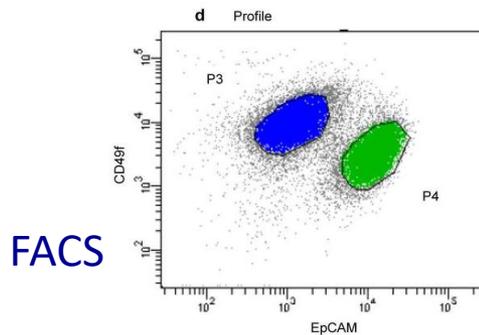
- Genomic



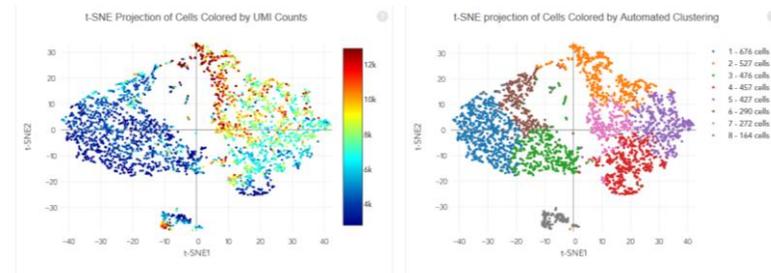
Transcription

Epigenetic

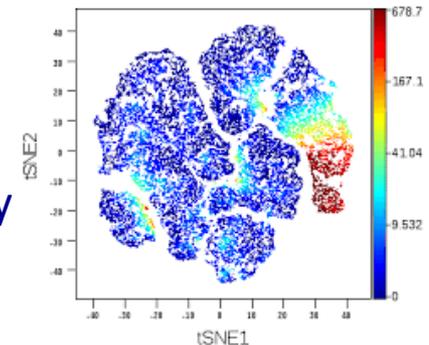
- Proteomic



## 10x scRNA-Seq

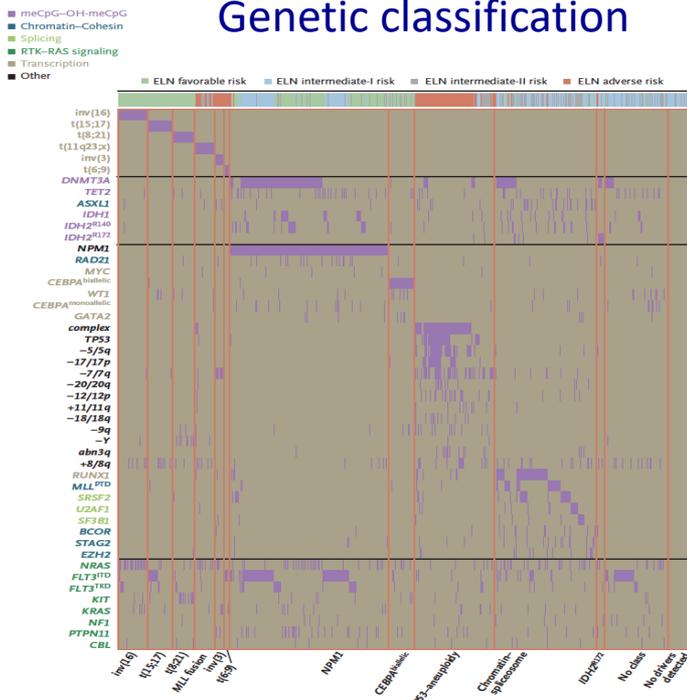


## CyTOF Mass Cytometry

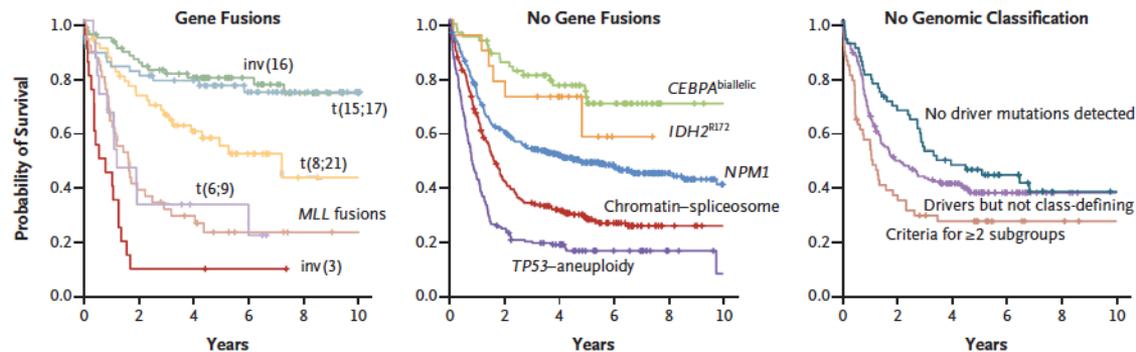


# Heterogeneity between tumours: the basis for tumour classification and disease prognostication

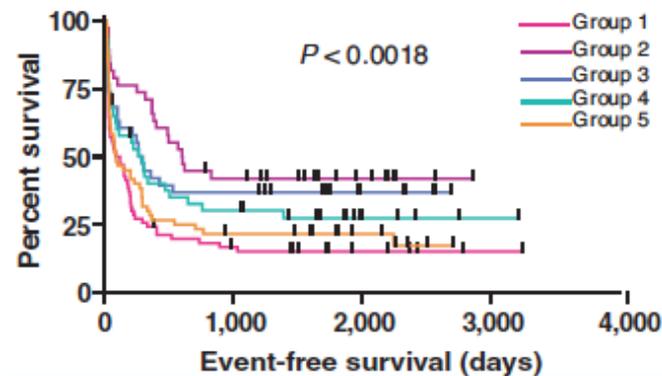
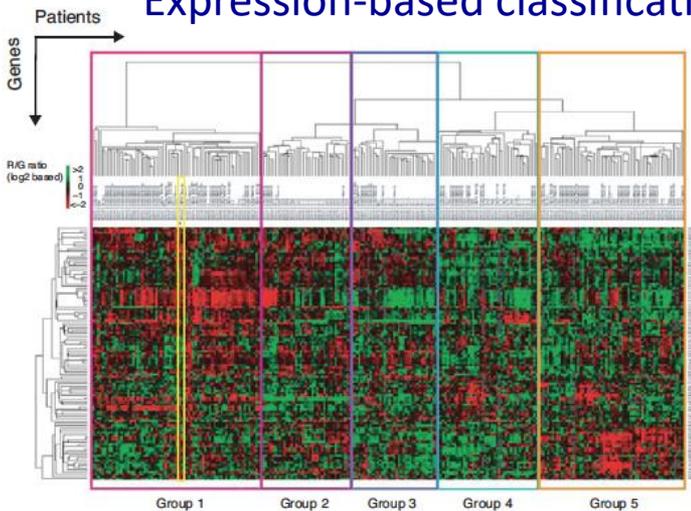
## Genetic classification



## OUTCOME MEASURES

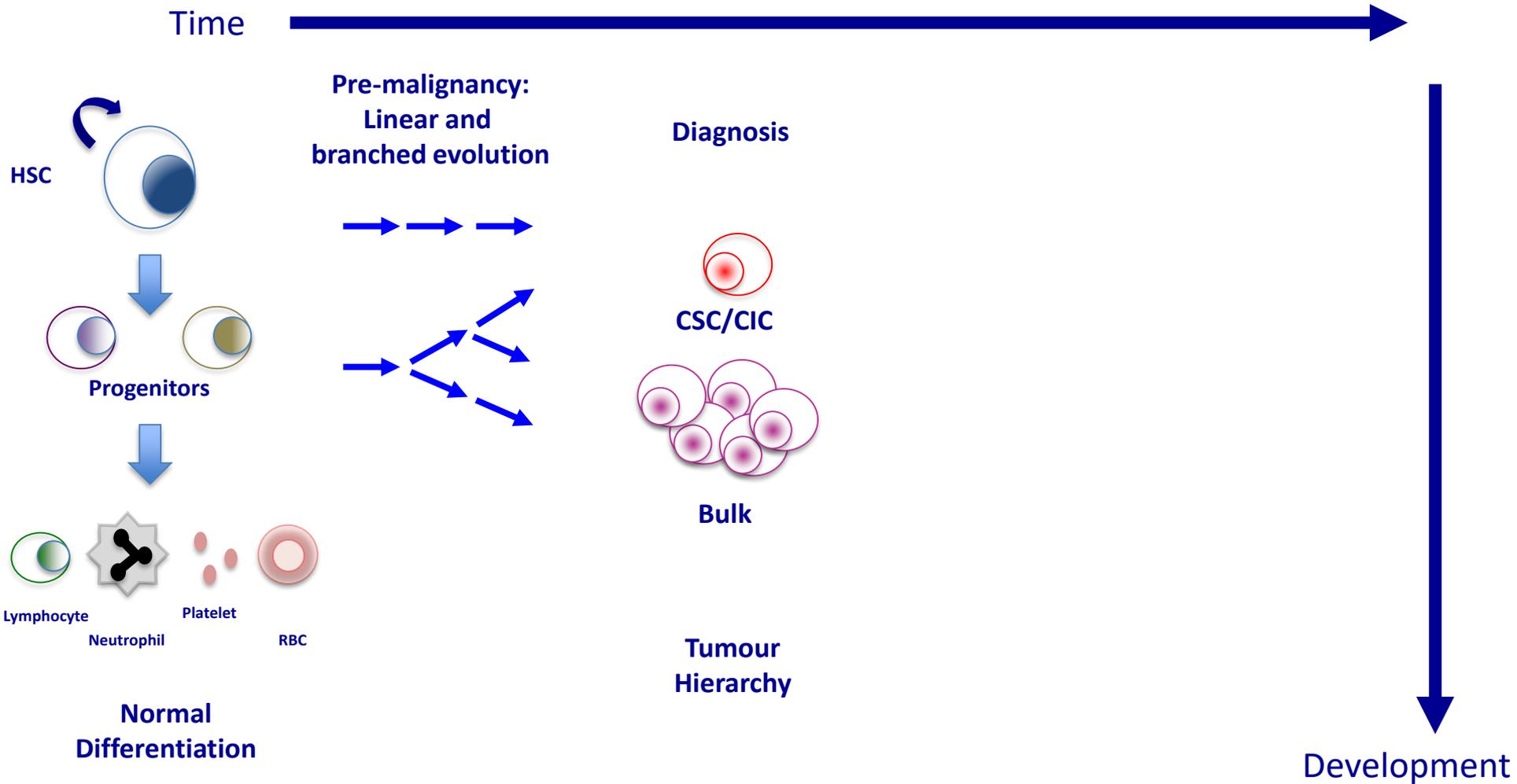


## Expression-based classification

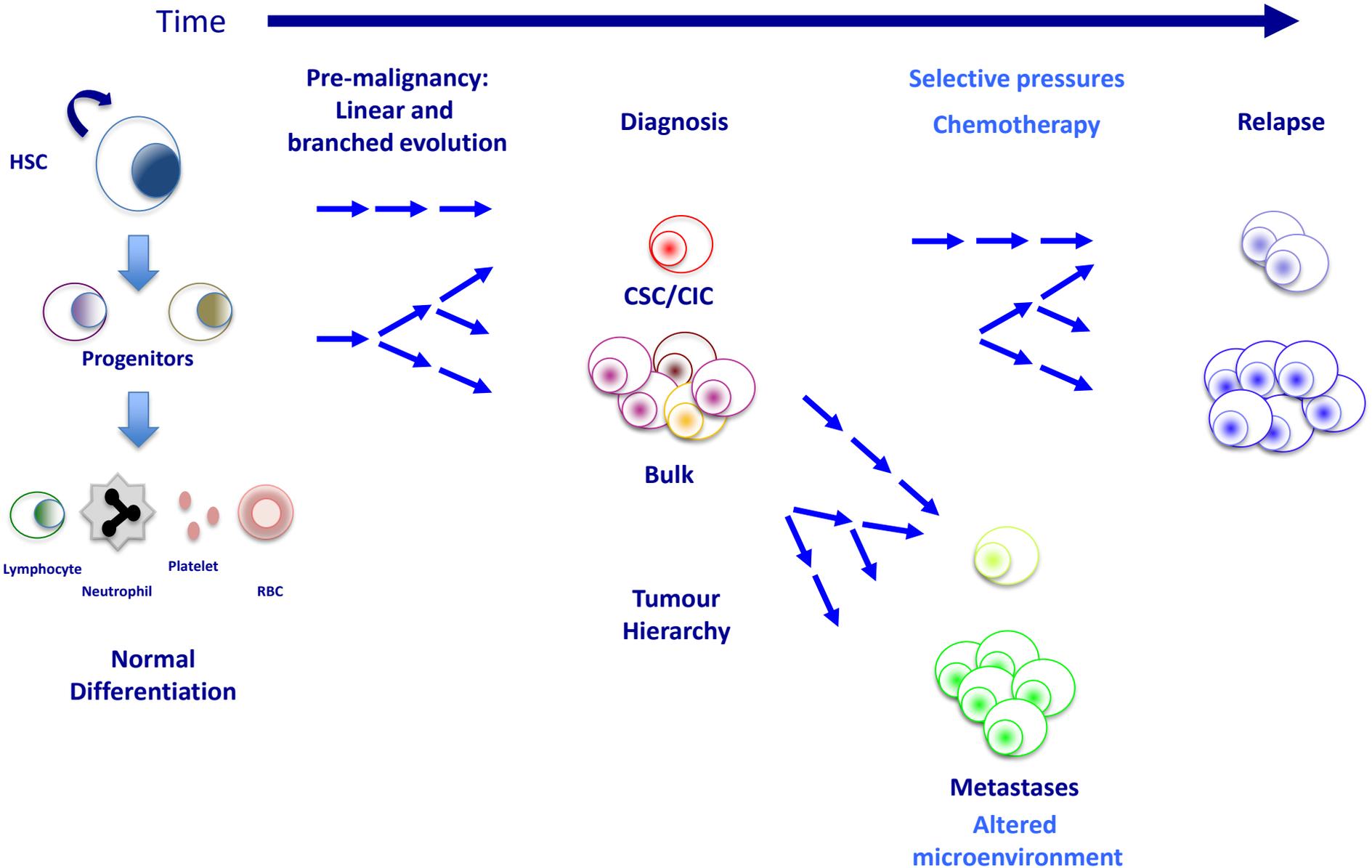


CLASSIFIERS

# How is heterogeneity generated within tumours?



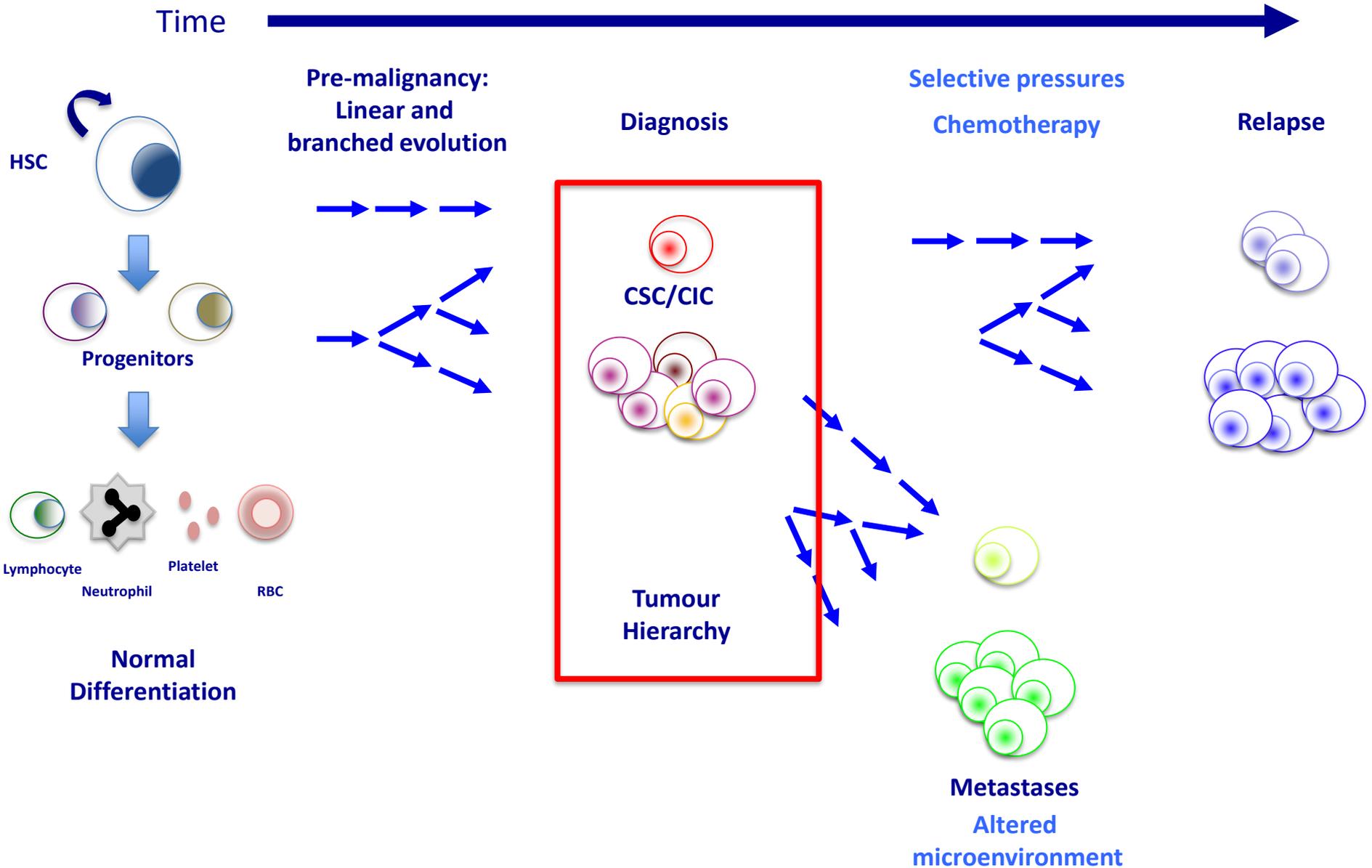
# Heterogeneity in Cancer- definition of terms



# Talk outline

- Context and definitions
- Tumour hierarchy: the cancer stem cell hypothesis and its implications for tumour biology
- Pre-malignancy – overt and covert premalignant states and early tumour evolution
- Tumour evolution under selective pressures:
  - Relapse
  - Metastases
- Clinical implications: Targeting Cancer stem cells  
Targeting tumour heterogeneity

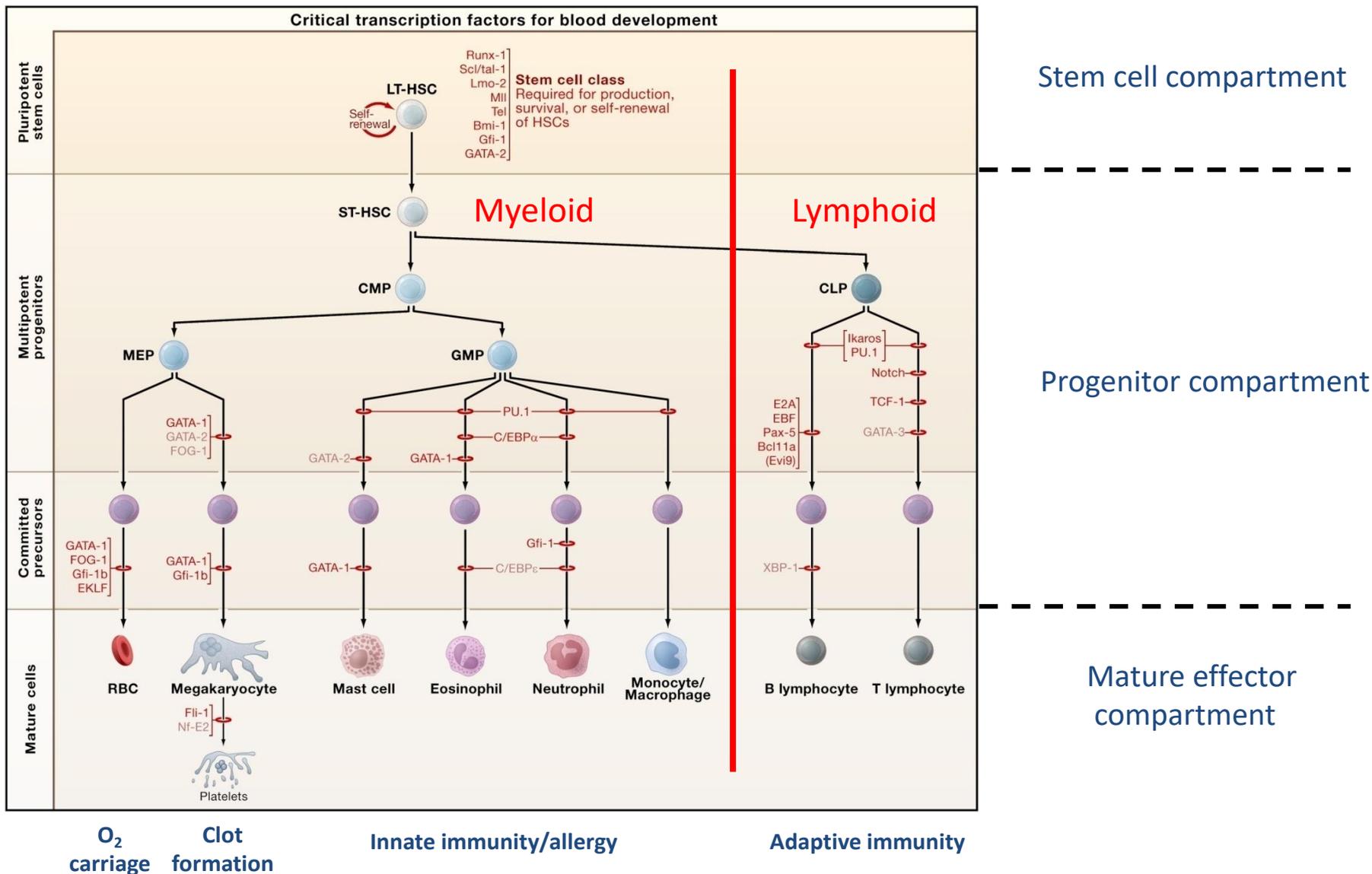
# Heterogeneity in Cancer- definition of terms



# The basis and implications of tumour heterogeneity

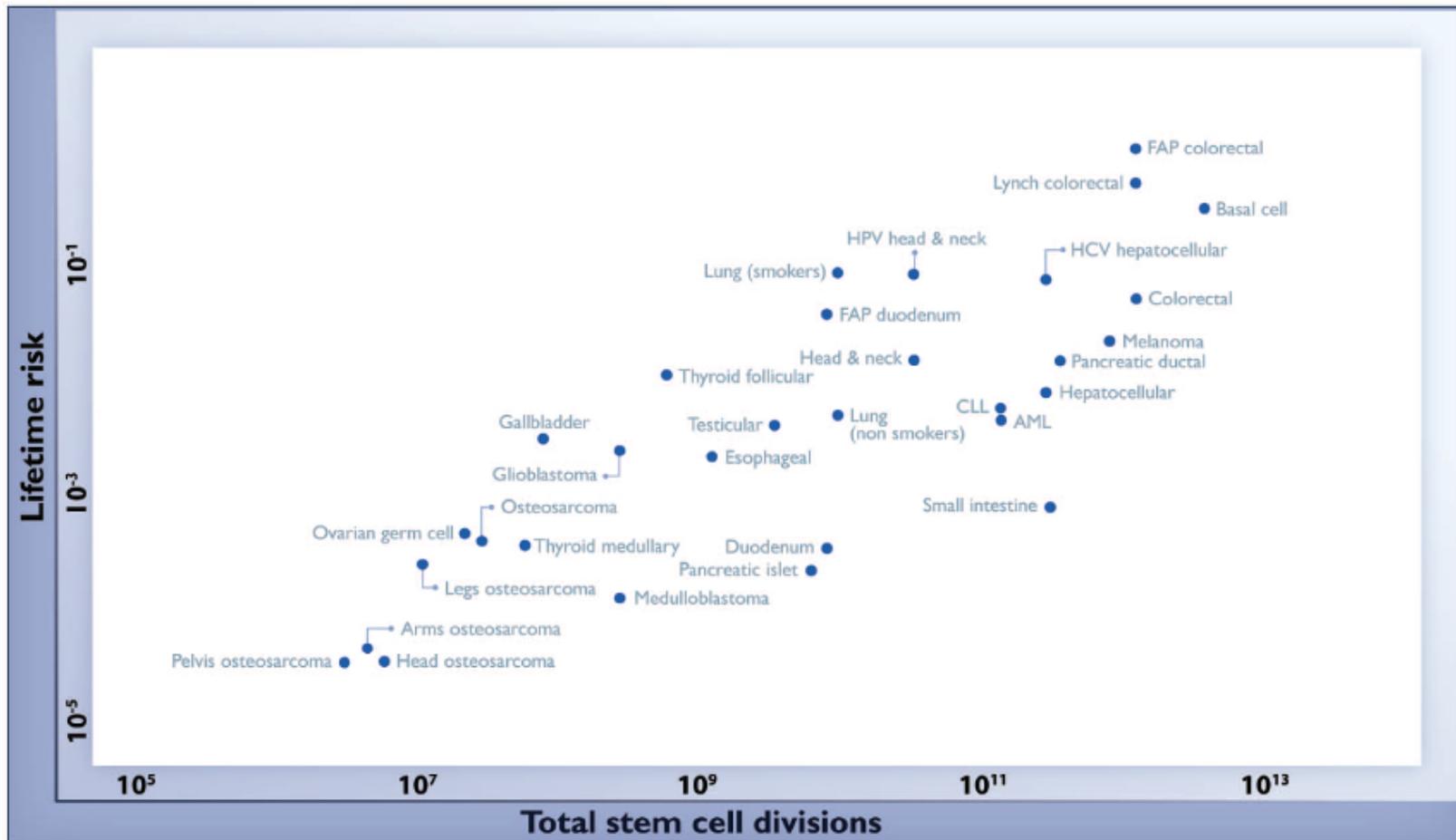
- Pathologists have noted intratumour heterogeneity visually for > century
- Advent of tumour assays and model systems – similar functional heterogeneity
- Till and McCulloch – suggest existence of haematopoietic stem cell (HSC)
- 1963 - Only 1-4% of murine lymphoma cells transplant into recipients
- 1965 - Similar for solid organ tumours in soft agar
- Autologous tumour inoculations into forearms of human subjects (!) – need  $> 1 \times 10^6$  cells to initiate a tumour
- ? Existence of stem cells in cancer

# Hierarchical organisation of adult mitotic organ systems



Cancers retain vestiges of normal tissue organisation

# Tissue cancer risk relates to stem cell frequency and rate of division?



FAP = Familial Adenomatous Polyposis ♦ HCV = Hepatitis C virus ♦ HPV = Human papillomavirus ♦ CLL = Chronic lymphocytic leukemia ♦ AML = Acute myeloid leukemia

Controversial –due to modeling?

Use of neutral mutations to model stem cell number and divisions

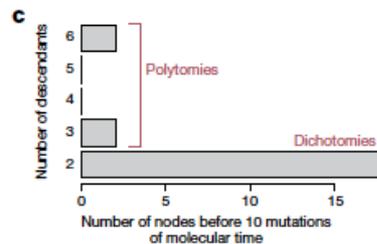
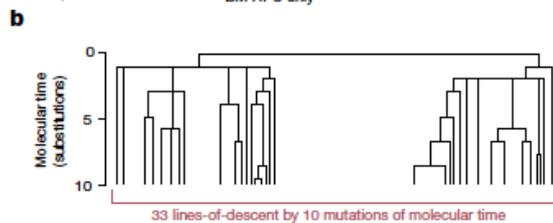
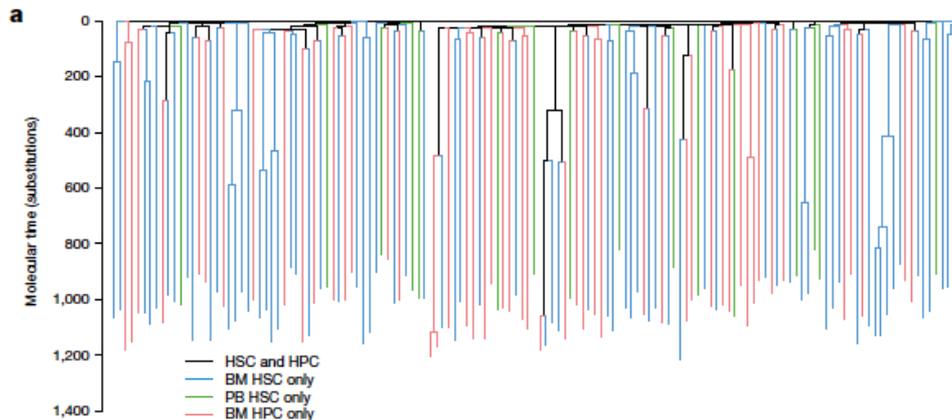
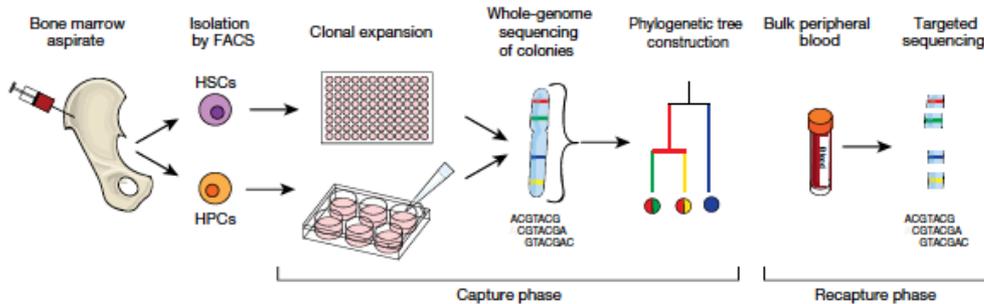
Tomasetti and Vogelstein, Science, 2015

# Use of neutral mutations to model stem cell number and dynamics

ARTICLE

## Population dynamics of normal human blood inferred from somatic mutations

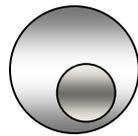
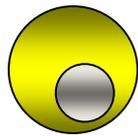
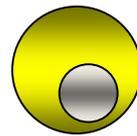
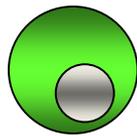
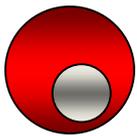
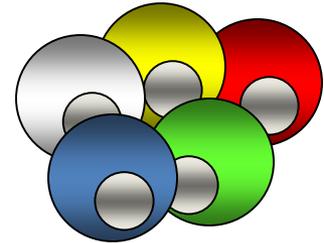
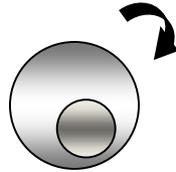
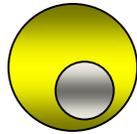
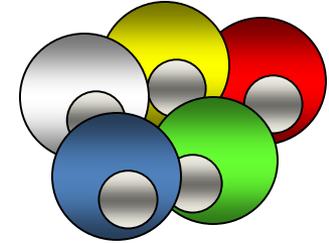
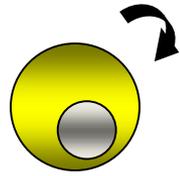
Henry Lee-Six<sup>1</sup>, Nina Friesgaard Øbro<sup>2</sup>, Mairi S. Shepherd<sup>2</sup>, Sebastian Grossmann<sup>1</sup>, Kevin Dawson<sup>1</sup>, Miriam Belmonte<sup>2</sup>, Robert J. Osborne<sup>3</sup>, Brian J. P. Huntly<sup>2</sup>, Inigo Martincorena<sup>1</sup>, Elizabeth Anderson<sup>1</sup>, Laura O'Neill<sup>1</sup>, Michael R. Stratton<sup>1</sup>, Elisa Laurenti<sup>2</sup>, Anthony R. Green<sup>2,3\*</sup>, David G. Kent<sup>2,3\*</sup> & Peter J. Campbell<sup>1,3\*</sup>



- Use neutral mutations to model stem cell number and dynamics
- Homeostatic measurement
- Human
- Suggests ~ 100,000 Haematopoietic stem cells (HSC)
- Range (44,000-215,000)
- Every HSC divides every 2-20 months

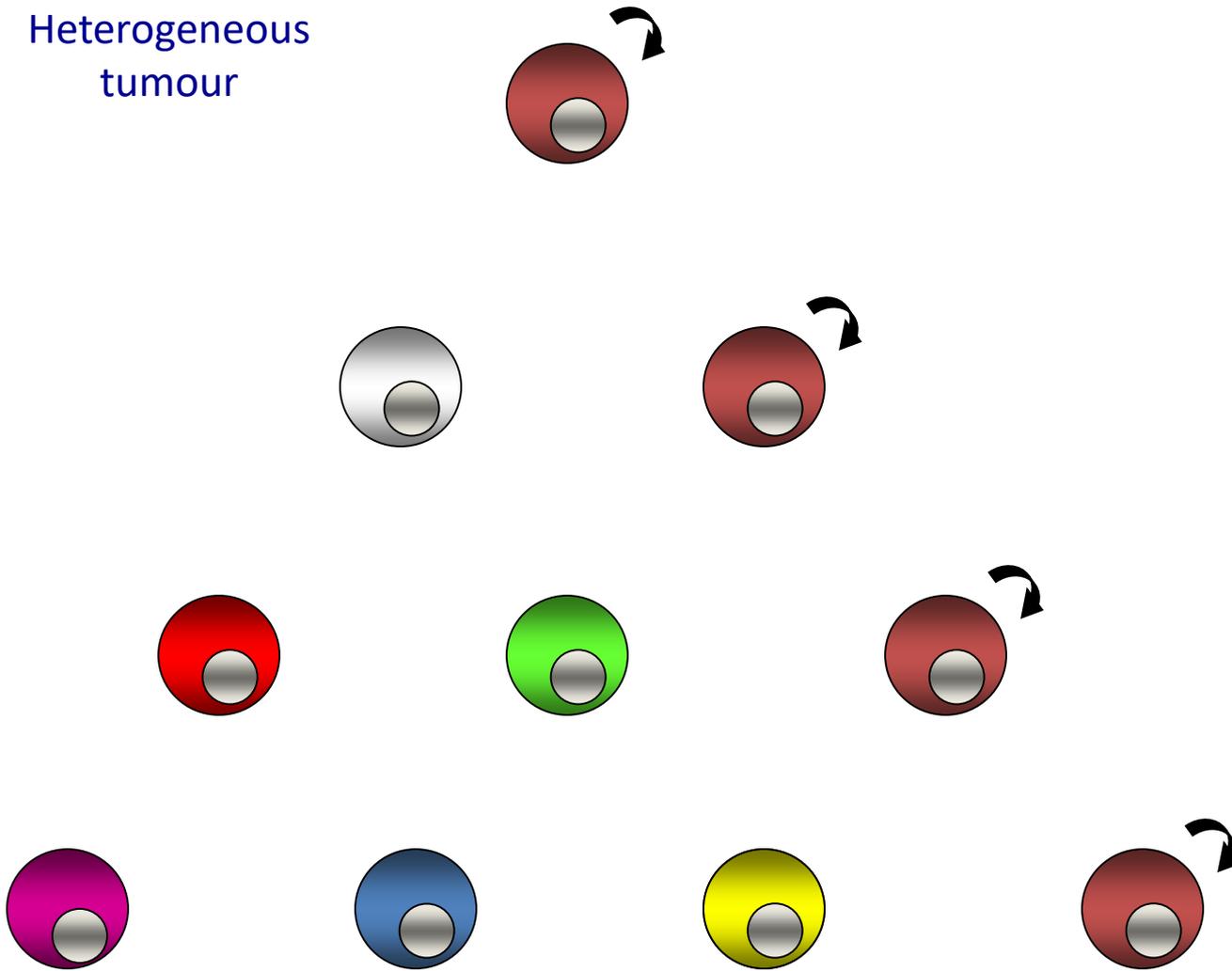
# Models of Cancer organisation and Heterogeneity: Stochastic/deterministic model

Heterogeneous  
tumour

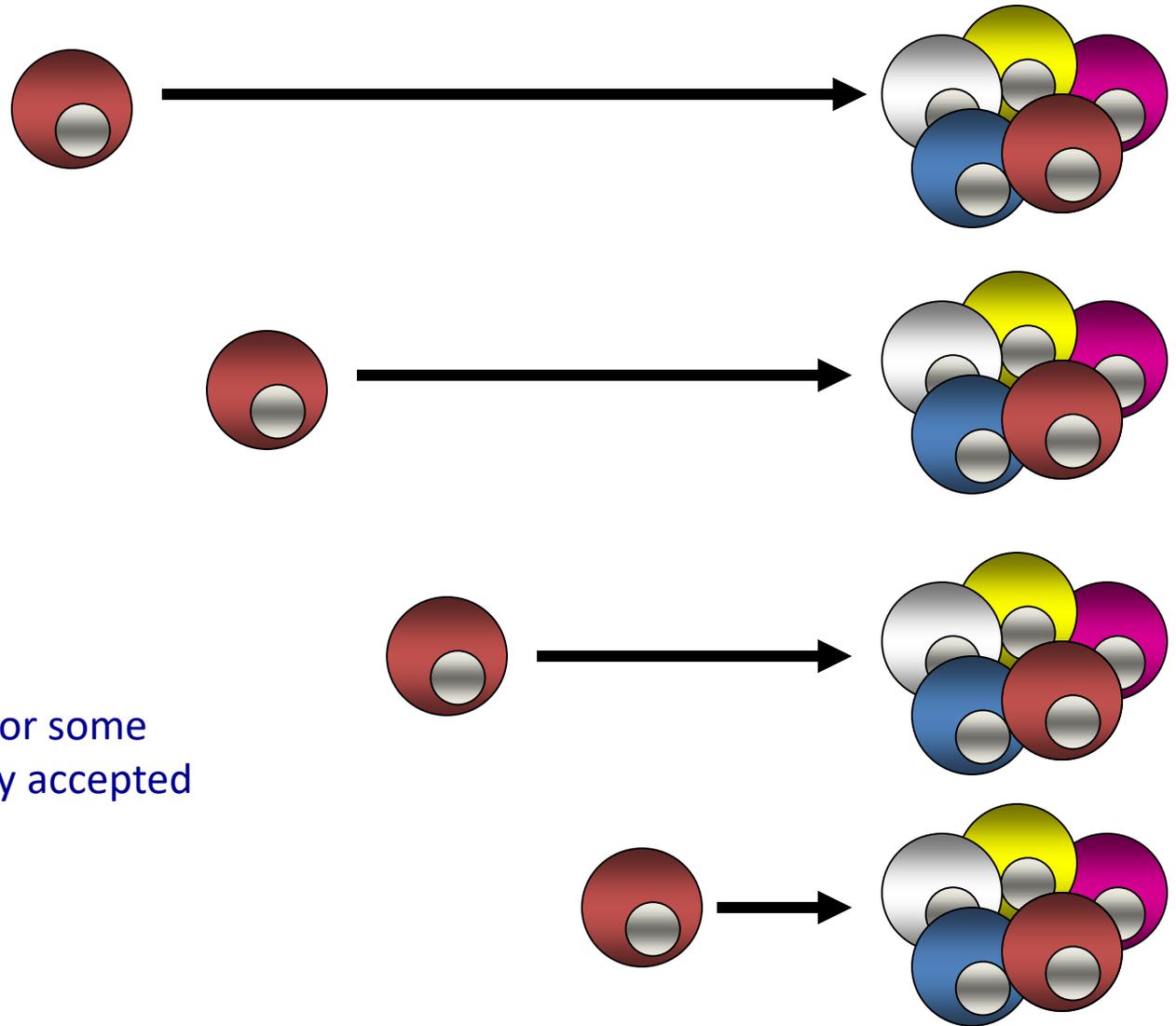


# Cancer is also arranged as a hierarchy: Cancer stem cell model

Heterogeneous  
tumour



# Cancer stem/initiating cell model



Concept is controversial for some malignancies but is widely accepted in most malignancies

# Demonstration of cancer (leukaemia) stem cells in human acute myeloid leukaemia (AML)

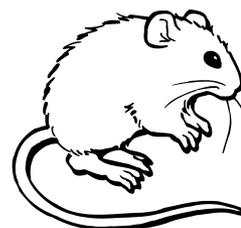
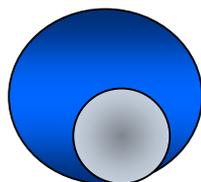
Human acute myeloid leukemia is organized as a hierarchy that originates from a primitive hematopoietic cell

DOMINIQUE BONNET & JOHN E. DICK

*Department of Genetics, Research Institute, Hospital for Sick Children and Department of Molecular and Medical Genetics, University of Toronto, 555 University Avenue, Toronto, Ontario M5G 1X8, Canada  
Correspondence should be addressed to J.E.D.*

*Department of Genetics, Research Institute, Hospital for Sick Children, 555 University Avenue, Toronto, Ontario, Canada M5G 1X8*

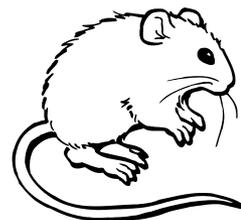
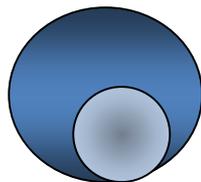
Immature CD34+/38-



LEUKAEMIA

Leukaemia  
cells  
from  
patients

More mature CD34+/38+



NO LEUKAEMIA

# Demonstration of cancer stem cells in other Solid organ malignancies

## Prospective identification of tumorigenic breast cancer cells

PNAS, 2003

Muhammad Al-Hajj\*, Max S. Wicha\*, Adalberto Benito-Hernandez<sup>†</sup>, Sean J. Morrison\*\*<sup>‡§</sup>, and Michael F. Clarke\*\*<sup>†¶</sup>

Departments of \*Internal Medicine and <sup>†</sup>Pathology, Comprehensive Cancer Center, <sup>‡</sup>Department of Developmental Biology, and <sup>§</sup>Howard Hughes Medical Institute, University of Michigan Medical School, Ann Arbor, MI 48109

## Identification of human brain tumour initiating cells

Nature, 2004

Sheila K. Singh<sup>1,2,3</sup>, Cynthia Hawkins<sup>1,4</sup>, Ian D. Clarke<sup>1,2</sup>,  
Jeremy A. Squire<sup>6</sup>, Jane Bayani<sup>6</sup>, Takuichiro Hide<sup>1,2</sup>, R. Mark Henkelman<sup>5</sup>,  
Michael D. Cusimano<sup>3,7</sup> & Peter B. Dirks<sup>1,2,3</sup>

## A human colon cancer cell capable of initiating tumour growth in immunodeficient mice

Nature, 2006

Catherine A. O'Brien<sup>1</sup>, Aaron Pollett<sup>2</sup>, Steven Gallinger<sup>3</sup> & John E. Dick<sup>1,4</sup>

## Identification and expansion of human colon-cancer-initiating cells

Nature, 2006

Lucia Ricci-Vitiani<sup>1</sup>, Dario G. Lombardi<sup>2</sup>, Emanuela Pilozzi<sup>3</sup>, Mauro Biffoni<sup>1</sup>, Matilde Todaro<sup>4</sup>, Cesare Peschle<sup>1</sup>  
& Ruggero De Maria<sup>1,2</sup>

## Identification of Pancreatic Cancer Stem Cells

Chenwei Li,<sup>1</sup> David G. Heidt,<sup>1</sup> Piero Dalerba,<sup>4</sup> Charles F. Burant,<sup>2,3</sup> Lanjing Zhang,<sup>3</sup>  
Volkan Adsay,<sup>4</sup> Max Wicha,<sup>3</sup> Michael F. Clarke,<sup>5</sup> and Diane M. Simeone<sup>1,2</sup>

Cancer Res, 2007

Departments of <sup>1</sup>Surgery, <sup>2</sup>Molecular and Integrative Physiology, and <sup>3</sup>Internal Medicine, University of Michigan Medical Center, Ann Arbor, Michigan; <sup>4</sup>Department of Pathology, Karmanos Cancer Center, Detroit, Michigan; and <sup>5</sup>Department of Internal Medicine, Stanford University School of Medicine, Palo Alto, California

# The Empire strikes back: evidence against Cancer Stem Cells

Vol 456 | 4 December 2008 | doi:10.1038/nature07567

nature

## ARTICLES

Nature 2008

---

### **Efficient tumour formation by single human melanoma cells**

Elsa Quintana<sup>1\*</sup>, Mark Shackleton<sup>1\*</sup>, Michael S. Sabel<sup>2</sup>, Douglas R. Fullen<sup>3</sup>, Timothy M. Johnson<sup>4</sup> & Sean J. Morrison<sup>1</sup>

### **Tumor Growth Need Not Be Driven by Rare Cancer Stem Cells**

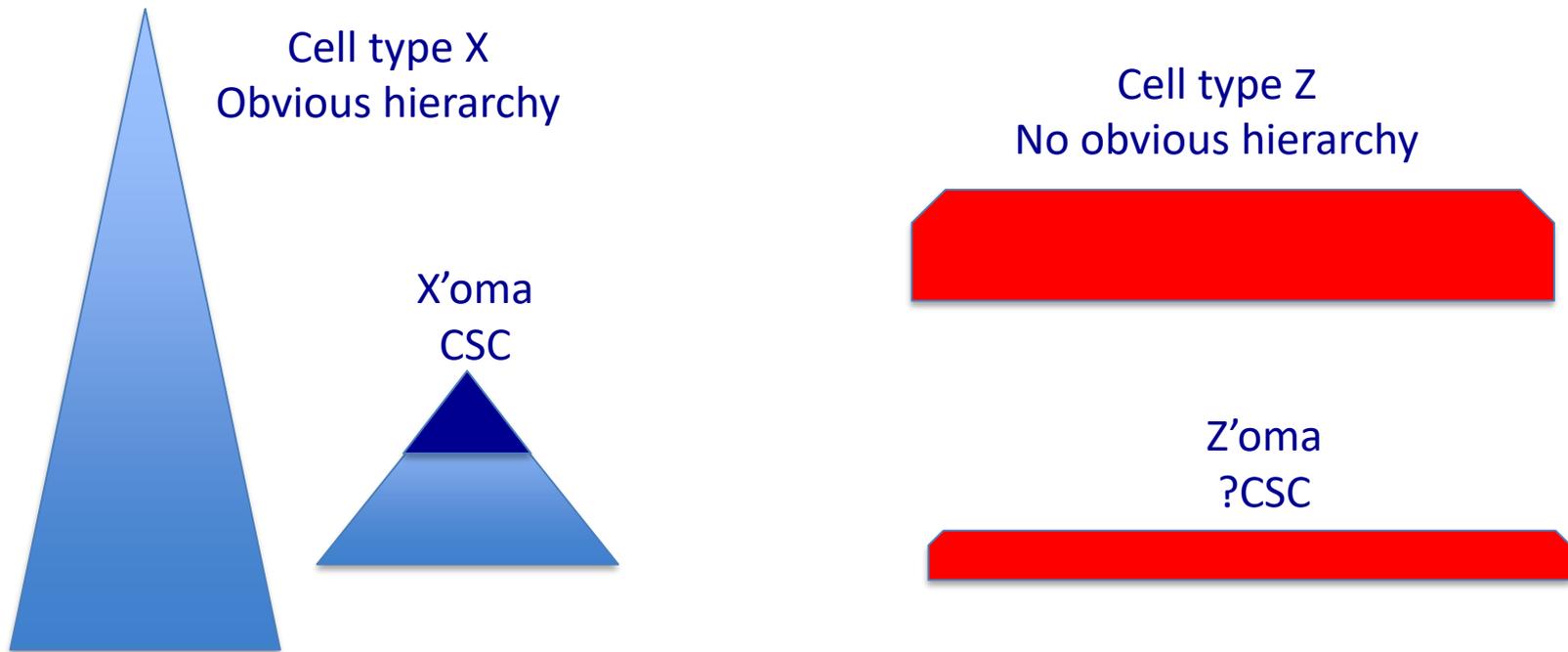
Science 2007

Priscilla N. Kelly,<sup>1,2</sup> Aleksandar Dakic,<sup>1,2</sup> Jerry M. Adams,<sup>1\*</sup>  
Stephen L. Nutt,<sup>1\*</sup> Andreas Strasser<sup>1\*†</sup>

- Similar assays
- High Frequency of cancer stem cells
- Lack of or shallow tissue Hierarchies

# How do we reconcile the evidence?

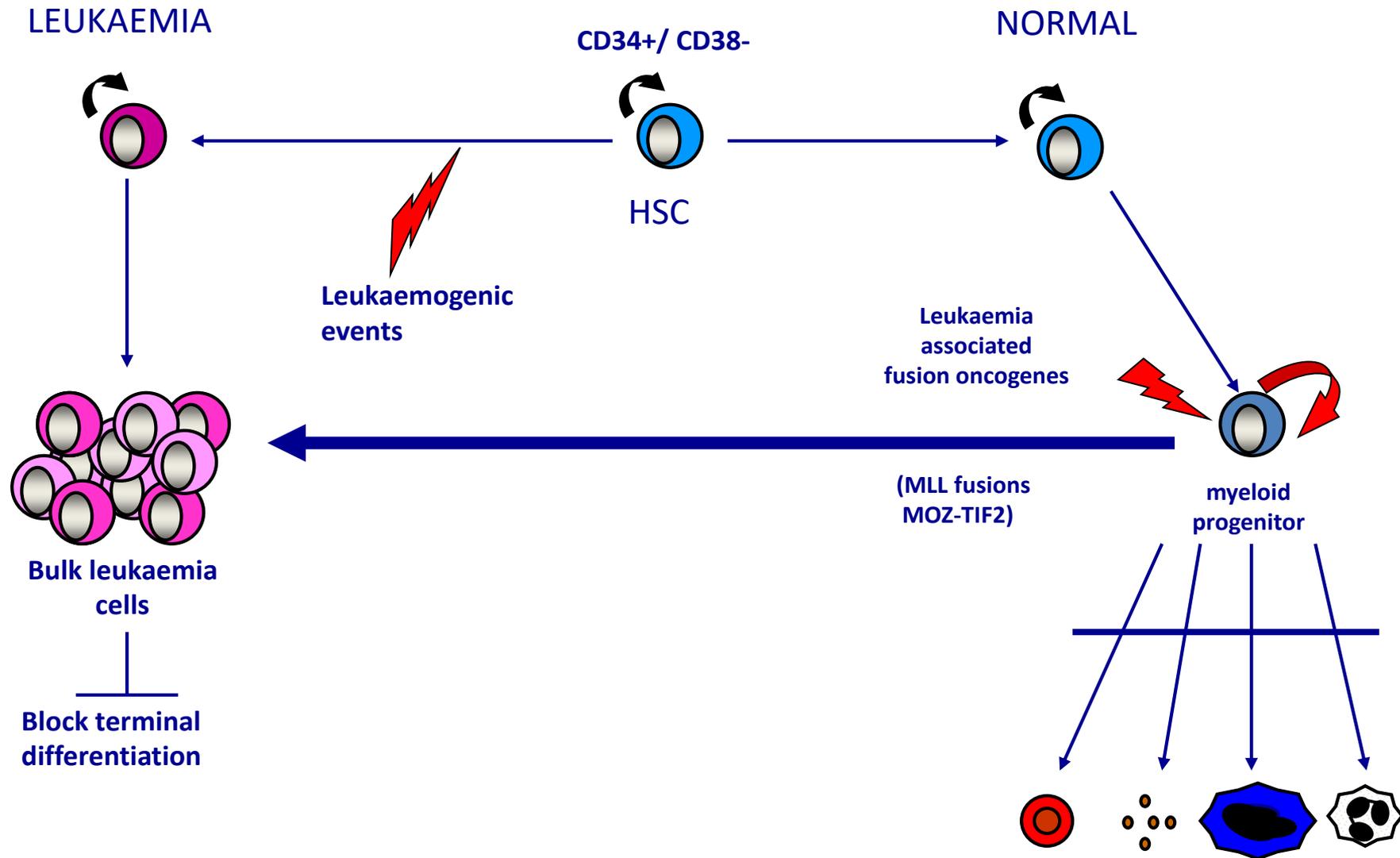
- One size does not fit all?
- Cancer stem cells do not need to be RARE populations
- Related to the hierarchy of the tissue from which tumour derived?



# Does the cell of origin determine heterogeneity

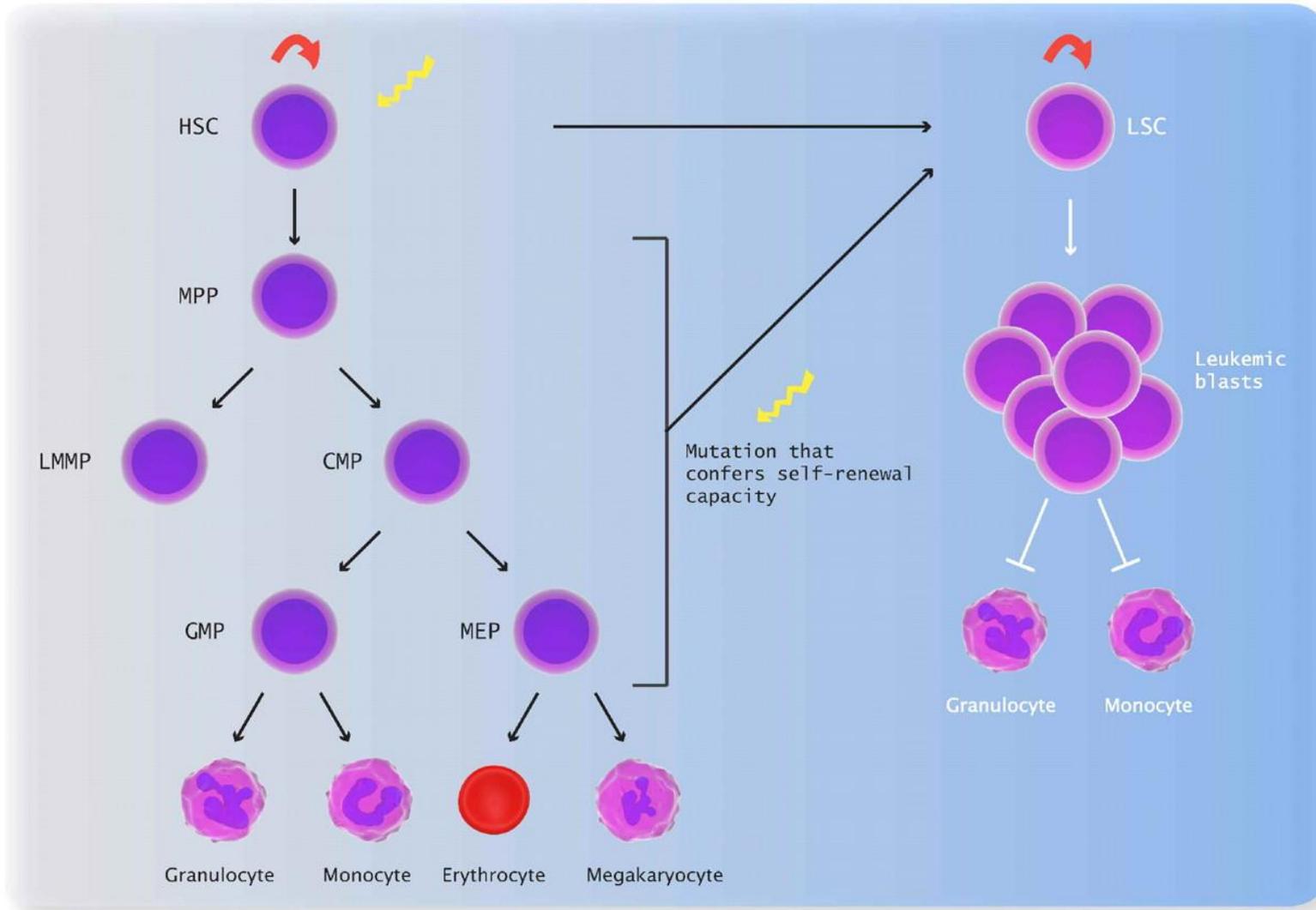
- Stem cell vs Non-stem cell: Benefits and limitations for both
- Stem cell:
  - Favourable – already has unlimited self-renewal, can accumulate the multiple mutations required for full malignant phenotype.
  - Unfavourable – Most important cell in metazoan biology, highly evolved DNA repair and DDR mechanisms to prevent mutation acquisition.  
Metabolic state.  
Quiescence > proliferation
- Progenitor/other cell:
  - Favourable- Genome less well protected.  
Metabolically more active  
Proliferation > Quiescence
  - Unfavourable – Very limited self-renewal – need to acquire this

# Committed progenitors may generate AML leukaemia stem cells



# AML leukaemia stem cells have surface characteristics of either stem or progenitor cells

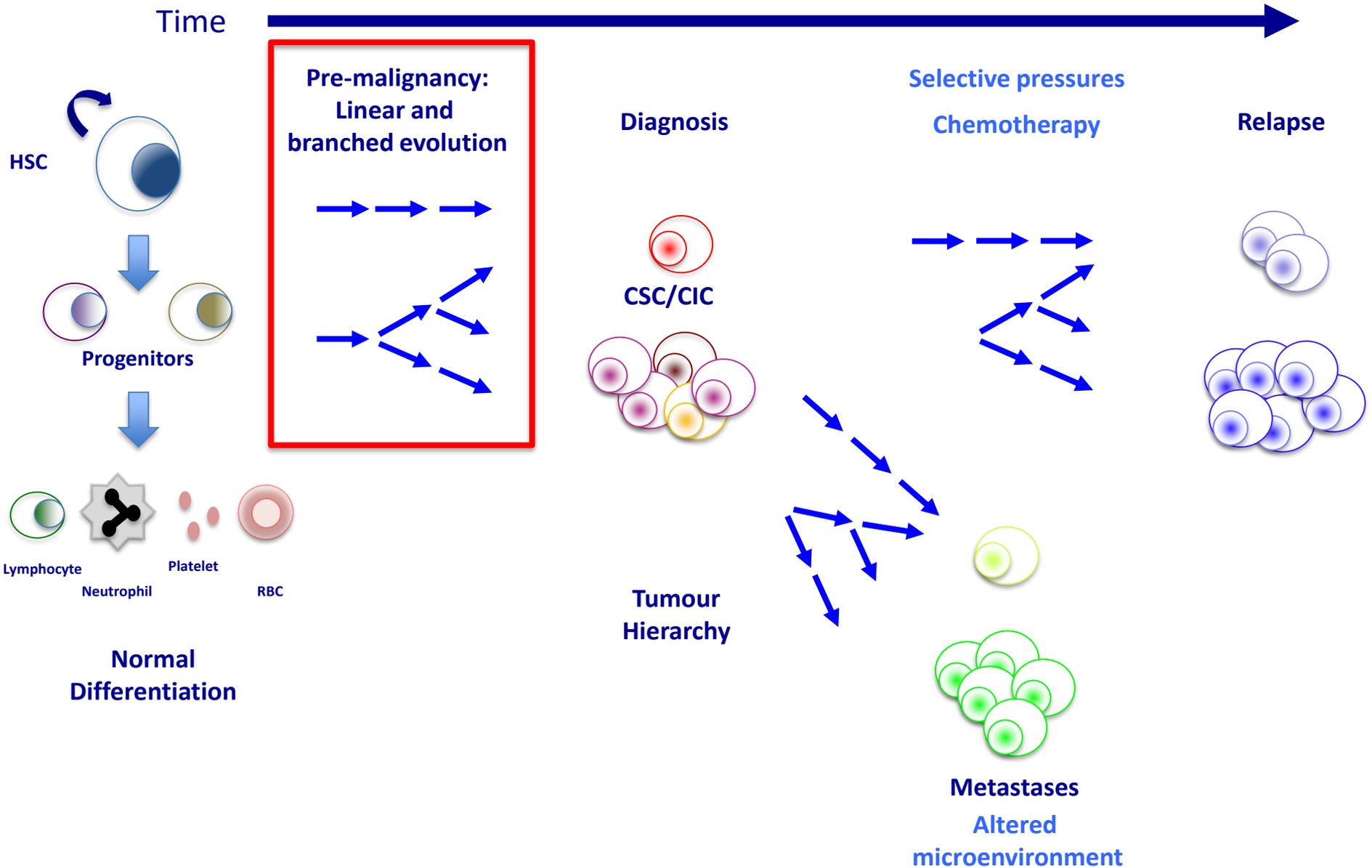
## Myeloid malignancies



# Cell of origin?

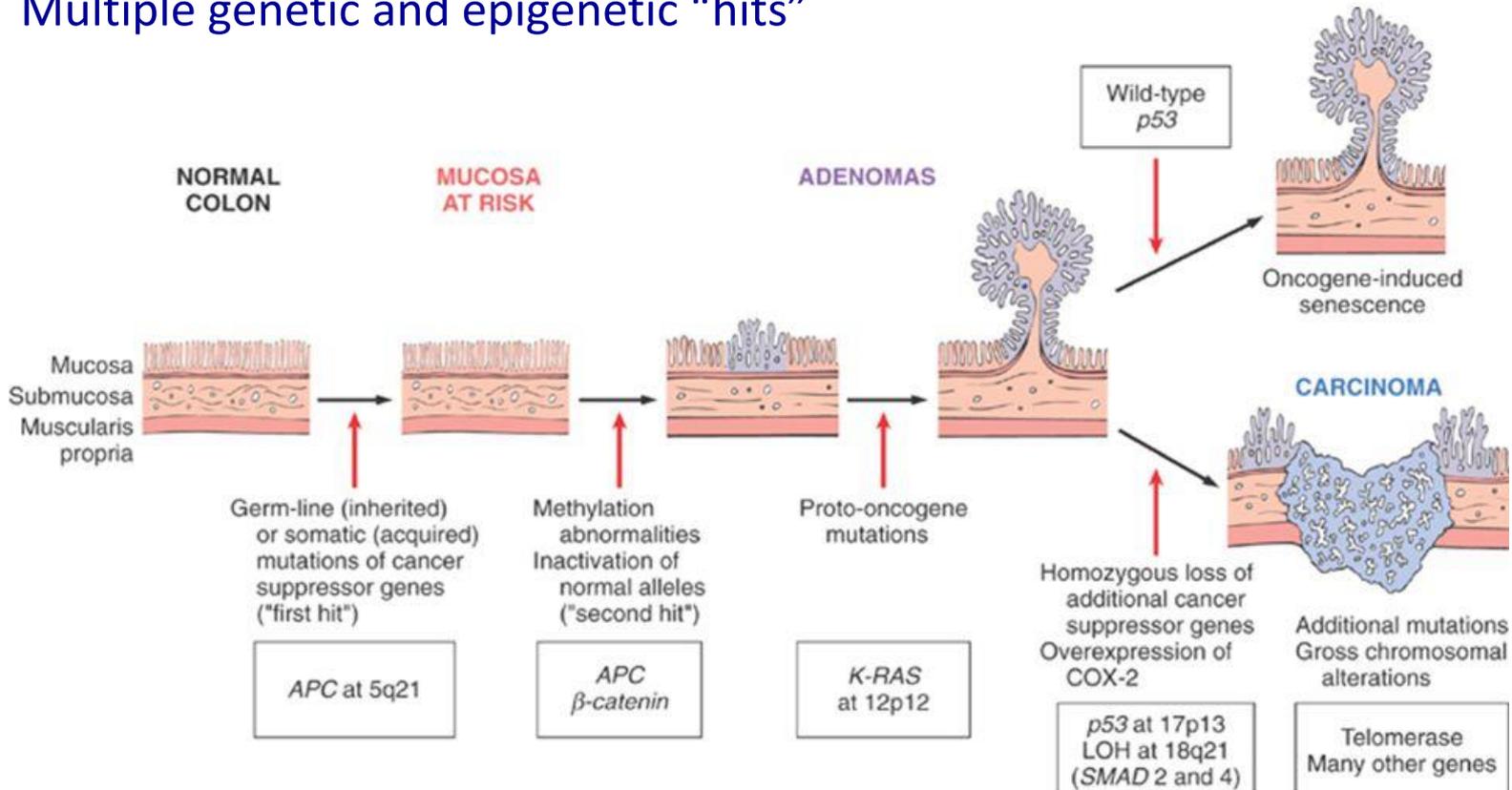
- Have to consider the stepwise acquisition of multiple mutations that is required to fully transform cells to the malignant state
- Some mutations as single events may alter progenitor cell function enough to generate a CSC
- Mutations may alter the frequency/probability of subsequent mutation acquisition
- Most likely that mutations are acquired in a stem cell but possible that subsequent downstream cells that acquire the full mutational complement are the tumour CSC

# Heterogeneity in Cancer: Pre-malignancy



# Premalignancy: Multi-step progression of cancer

Classical system – adenocarcinoma of the bowel  
 Multiple genetic and epigenetic “hits”



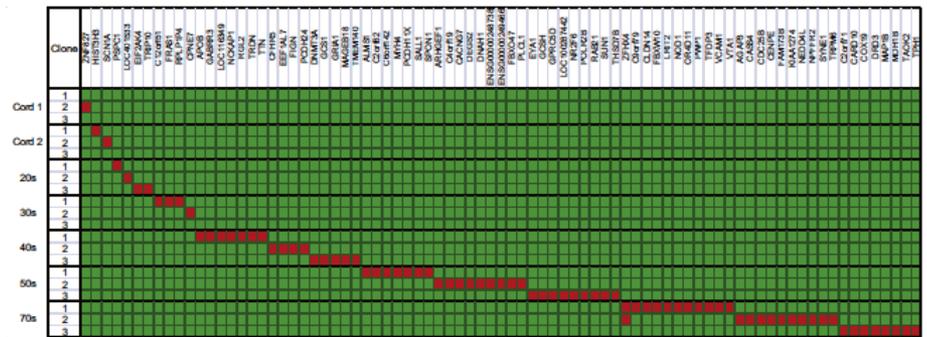
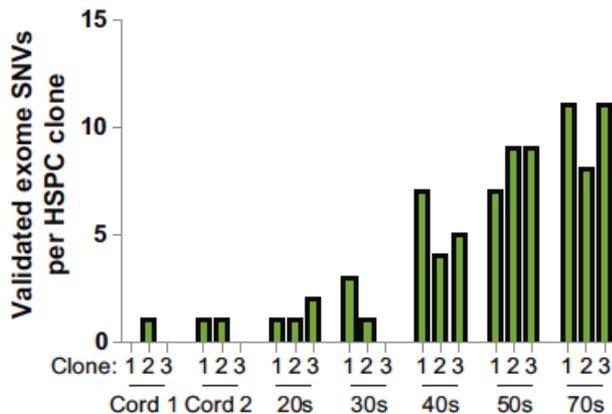
Overt phenotypes – dysplasia )  
 adenomatous polyps ) *in situ*

Screening in some individuals (FAP, Lynch syndrome) – surveillance and eradication

Similar for many other epithelial malignancies – Prostate, Pancreas, etc.

# Random mutation and Clonal Haematopoiesis

- Rate of mutation is one in every 50 million nucleotides (but  $\sim 6 \times 10^9$  bases in human genome!)
- Mutations exist in haematopoietic stem cells (HSC) from entirely normal individuals
- Function of age – may help to explain the later onset of most myeloid malignancies



Welch et al, Cell 2012

- Welch study - random mutations
- Suggests that normal HSC acquire random mutations as a function of time
- ?Stochastic whether they develop functional consequences/malignancy

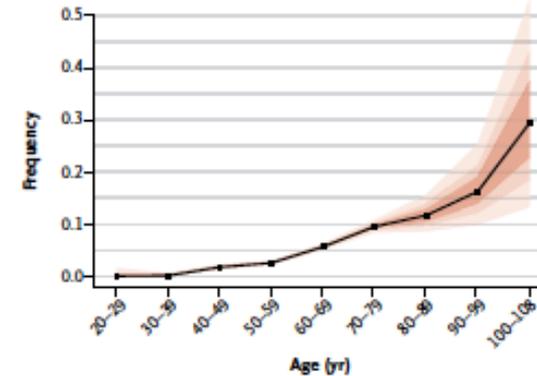
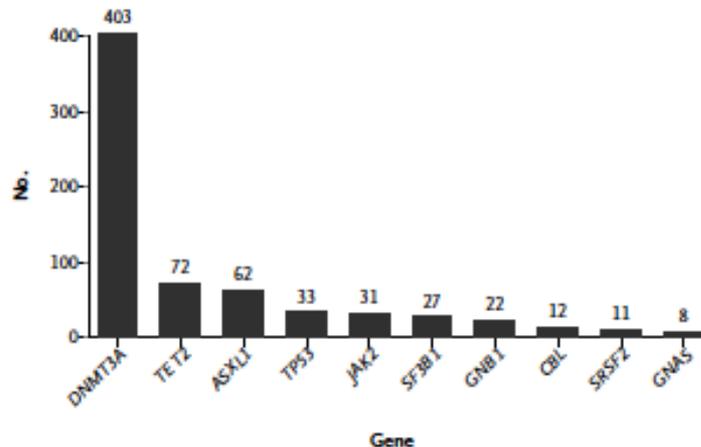
# Age related clonal Haematopoiesis (ARCH) Clonal Hameatopoiesis of indeterminate prognosis (CHIP)

The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

## Age-Related Clonal Hematopoiesis Associated with Adverse Outcomes

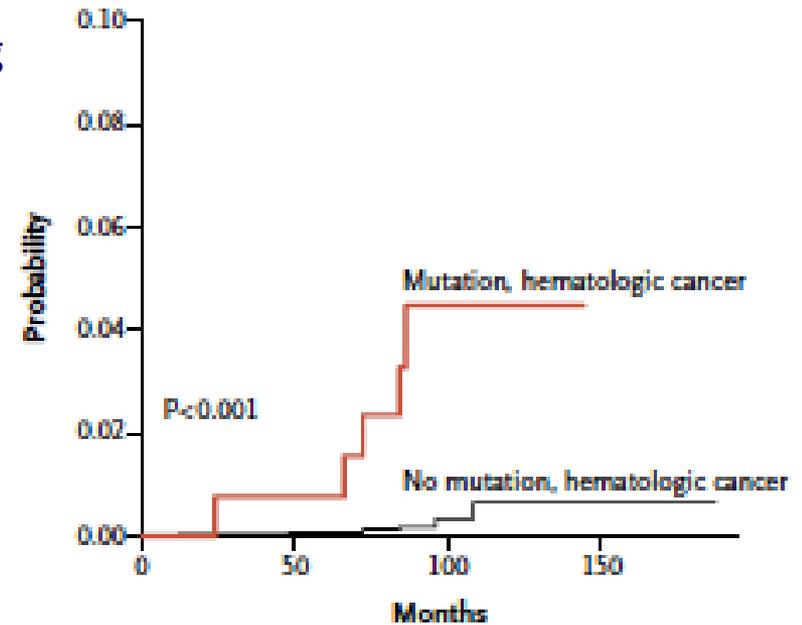
- WES data from 17,182 patients - mostly Type 2 diabetics for GWAS
- 805 somatic mutations in 73 genes from 746 individuals (4.5%) – age dependent phenomenon – rare under 40 but > 90 (18.4%).
- Majority in 3 genes - DNMT3A (403 VARIANTS) , TET2 (72) and ASXL1 (62)
- Median variant allelic frequency (VAF) 0.09 (i.e. 18%)



No. with Mutation	0	1	50	138	282	219	37	14	5
Total	240	855	2894	5441	5002	2300	317	86	17

# Age related clonal Haematopoiesis (ARCH)

- Median FU 8 years –
- Mutation associated with a HR of 11 of developing a haematological malignancy (95% CR 3.9-33)
- Assoc with an overall decreased survival (HR 1.4 and INDEPENDENT OF HAEM MALIGNANCY)
- NIDDM – OR 1.3
- IHD – OR 2.0
- CVA – OR 2.6



# Age Related Clonal Haematopoiesis (ARCH)

The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

## Clonal Hematopoiesis and Blood-Cancer Risk Inferred from Blood DNA Sequence

Age-related mutations associated with clonal hematopoietic expansion and malignancies

nature  
medicine

## Cell Reports

Report

## Leukemia-Associated Somatic Mutations Drive Distinct Patterns of Age-Related Clonal Hemopoiesis

EVIDENCE OF SIMILAR PREMALIGNANT PHENOTYPES THAT INCREASE WITH AGE IN OTHER TISSUES

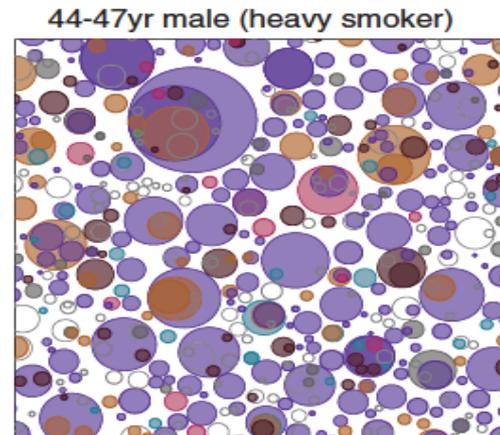
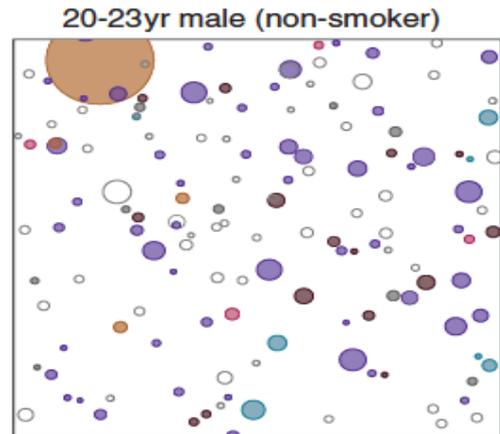
# Similar mutant clonal in epithelial tissues

## Somatic mutant clones colonize the human esophagus with age

Iñigo Martincorena<sup>1\*†</sup>, Joanna C. Fowler<sup>1\*</sup>, Agnieszka Wabik<sup>1</sup>, Andrew R. J. Lawson<sup>1</sup>, Federico Abascal<sup>1</sup>, Michael W. J. Hall<sup>1,2</sup>, Alex Cagan<sup>1</sup>, Kasumi Murai<sup>1</sup>, Krishnaa Mahbubani<sup>3</sup>, Michael R. Stratton<sup>1</sup>, Rebecca C. Fitzgerald<sup>2</sup>, Penny A. Handford<sup>4</sup>, Peter J. Campbell<sup>1,5</sup>, Kourosh Saeb-Parsy<sup>3</sup>, Philip H. Jones<sup>1†</sup>

9 cadaveric organ transplant donors

- Multiple cancer associated mutations found
- Increase frequency with age
- *NOTCH1* 12-80% and *TP53* 2-37%
- *NOTCH1* much higher in normal vs cancer



# Clonal evolution: Presence of pre-leukemia stem cells in AML

ARTICLE

doi:10.1038/nature13038

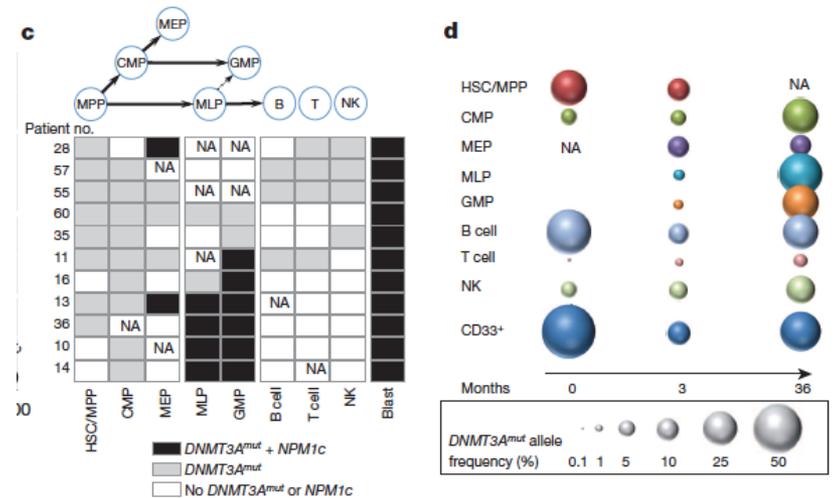
## Identification of pre-leukaemic haematopoietic stem cells in acute leukaemia

Liran I. Shlush<sup>1\*</sup>, Sasan Zandi<sup>1\*</sup>, Amanda Mitchell<sup>1</sup>, Weihsu Claire Chen<sup>1</sup>, Joseph M. Brandwein<sup>1,2,3</sup>, Vikas Gupta<sup>1,2,3</sup>, James A. Kennedy<sup>1</sup>, Aaron D. Schimmer<sup>1,2,3,4</sup>, Andre C. Schuh<sup>1,2,3</sup>, Karen W. Yee<sup>1,2,3</sup>, Jessica L. McLeod<sup>1</sup>, Monica Doedens<sup>1</sup>, Jessie J. F. Medeiros<sup>1</sup>, Rene Marke<sup>1,5</sup>, Hyeoung Joon Kim<sup>6</sup>, Kwon Lee<sup>6</sup>, John D. McPherson<sup>4,7</sup>, Thomas J. Hudson<sup>4,7,8</sup>, The HALT Pan-Leukemia Gene Panel Consortium†, Andrew M. K. Brown<sup>7</sup>, Fouad Yousif<sup>7</sup>, Quang M. Trinh<sup>7</sup>, Lincoln D. Stein<sup>7,8</sup>, Mark D. Minden<sup>1,2,3,4</sup>, Jean C. Y. Wang<sup>1,2,3</sup> & John E. Dick<sup>1,8</sup>

Nature 2014

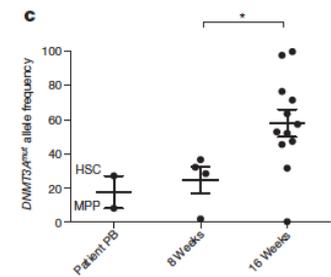
# Existence of Pre-leukaemia stem cells: Resistance to therapy and biological properties

- AML patient samples vs cultured T-cells – targeted sequencing 103 commonly mutated genes - 12 samples, 4 with *DNMT3A* mutation- ALSO PRESENT in 3/4 T-CELLS
- NOT contamination as co-occurring mutations NOT present (*NPM1c*)
- Present in normal differentiating haem tissue
- Preleukaemic stem cell with transformation of a lower cell population
- DNMT3A<sup>mut</sup>* pre-LSC survive chemotherapy
- DNMT3A<sup>mut</sup>* pre-LSC outcompete normal HSC and undergo clonal expansion in NSG mice

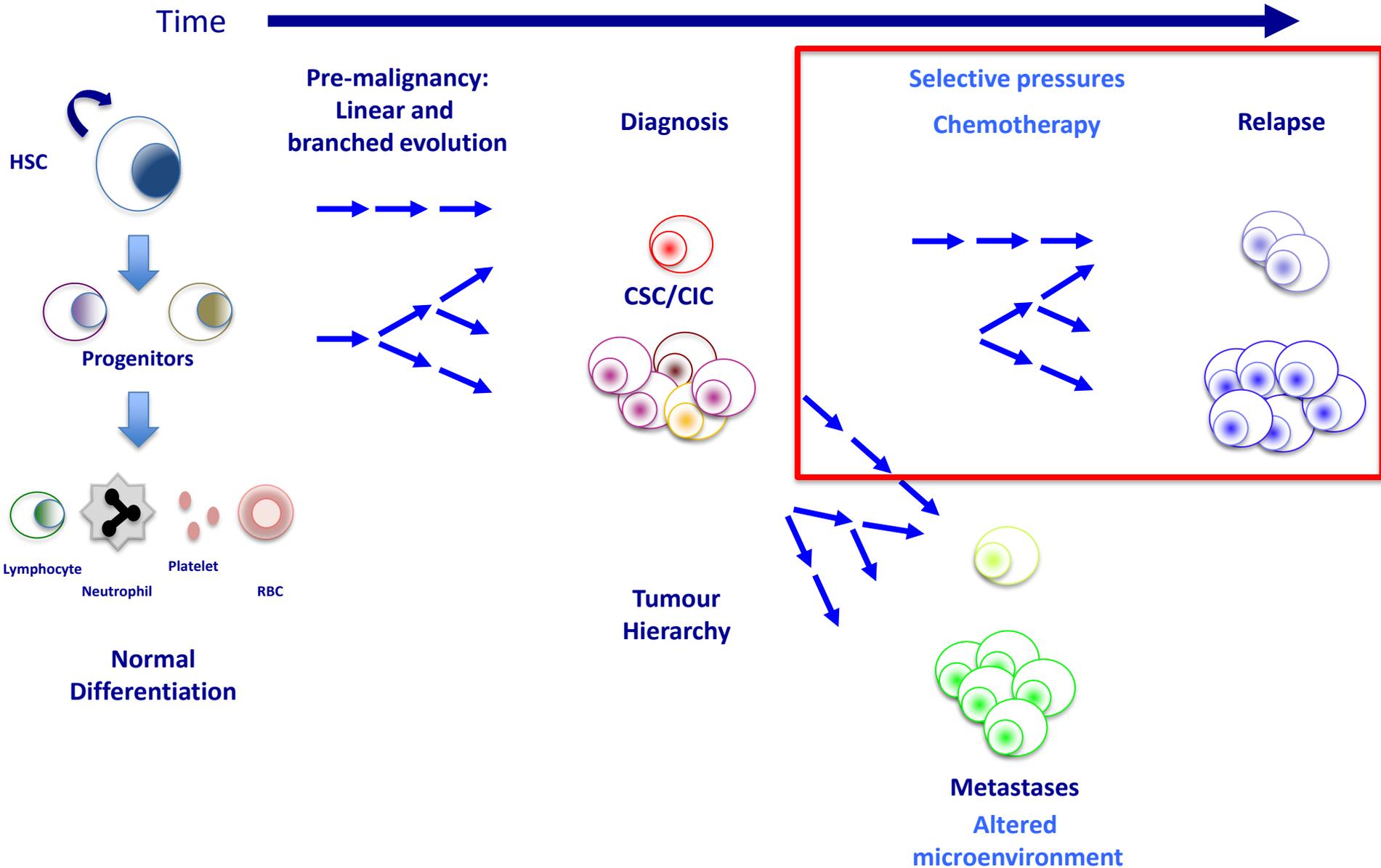


**b**

Mouse no.	Cell dose ( $\times 10^6$ )	Graft composition		Allele frequency	
		CD19 <sup>+</sup>	CD33 <sup>+</sup>	<i>DNMT3A<sup>mut</sup></i>	<i>NPM1c</i>
9	1.0	90.8	3.2		
2	2.0	80.2	6.5		
5	2.0	86.4	2.7		
25	0.5	92.1	0.8		
24	0.5	91.3	0.0		
4	2.0	85.4	4.6		
3	2.0	67.4	22.4		
12	1.0	41.5	53.2		
20	0.5	93.5	1.8		
11	1.0	78.7	7.1		
14	1.0	87.2	6.7		
10	1.0	89.8	1.8		



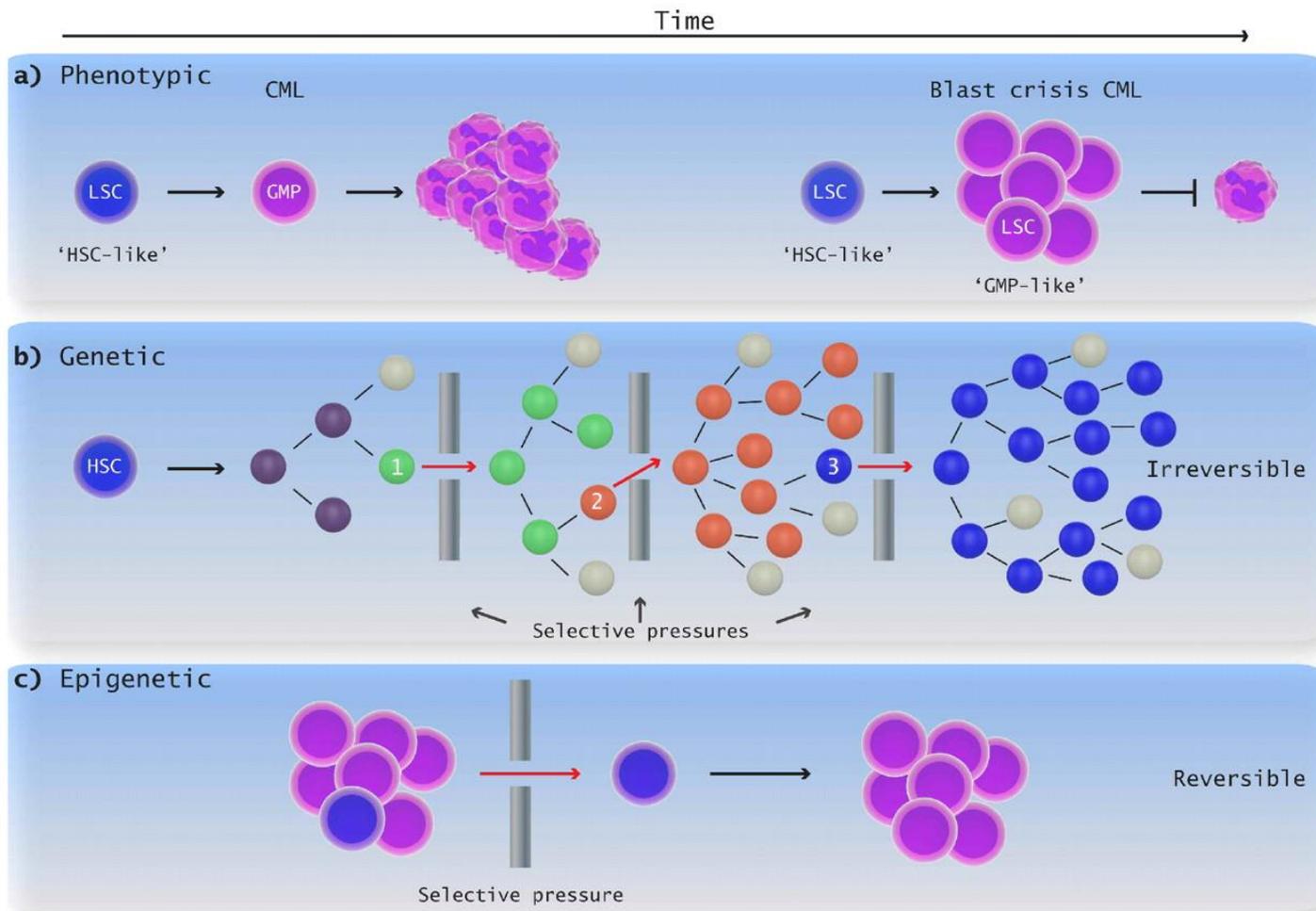
# Heterogeneity in Cancer- definition of terms



# Fatal consequences of cancer: Relapse

- We are actually pretty successful at treating primary cancers
- Surgery
- Radiotherapy
- Chemotherapy
- Immunotherapy
- NOT at preventing spread – Metastasis
- NOT at preventing recurrence – Relapse
- **Most patients die from relapse and/or spread of disease**

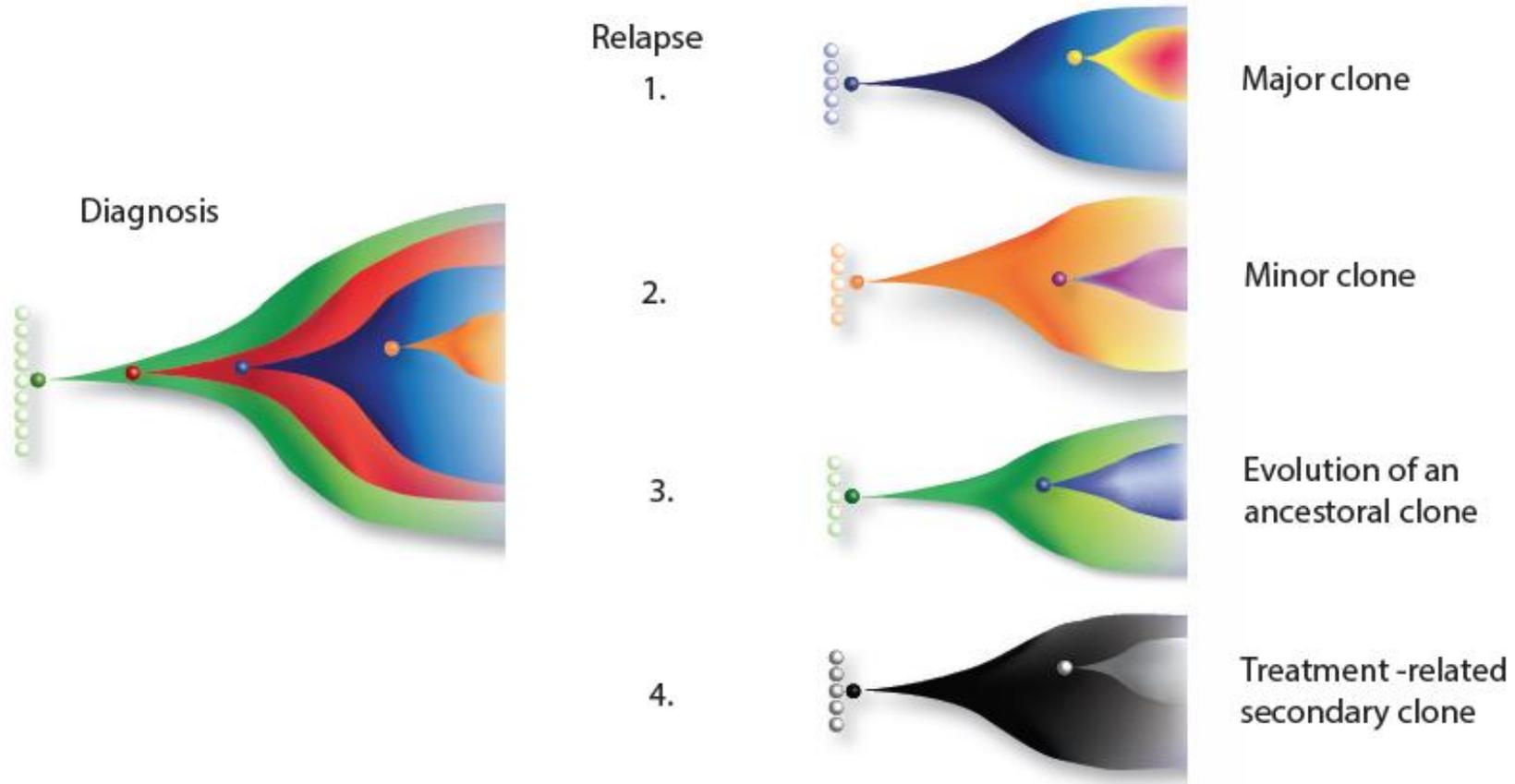
# Heterogeneity of Cancer Stem Cells: Evolution under selective pressures



- Tumours are constantly evolving under fixed and mobile selective pressures
- Therapy: standard chemo, DRT, immunotherapy and targeted therapy

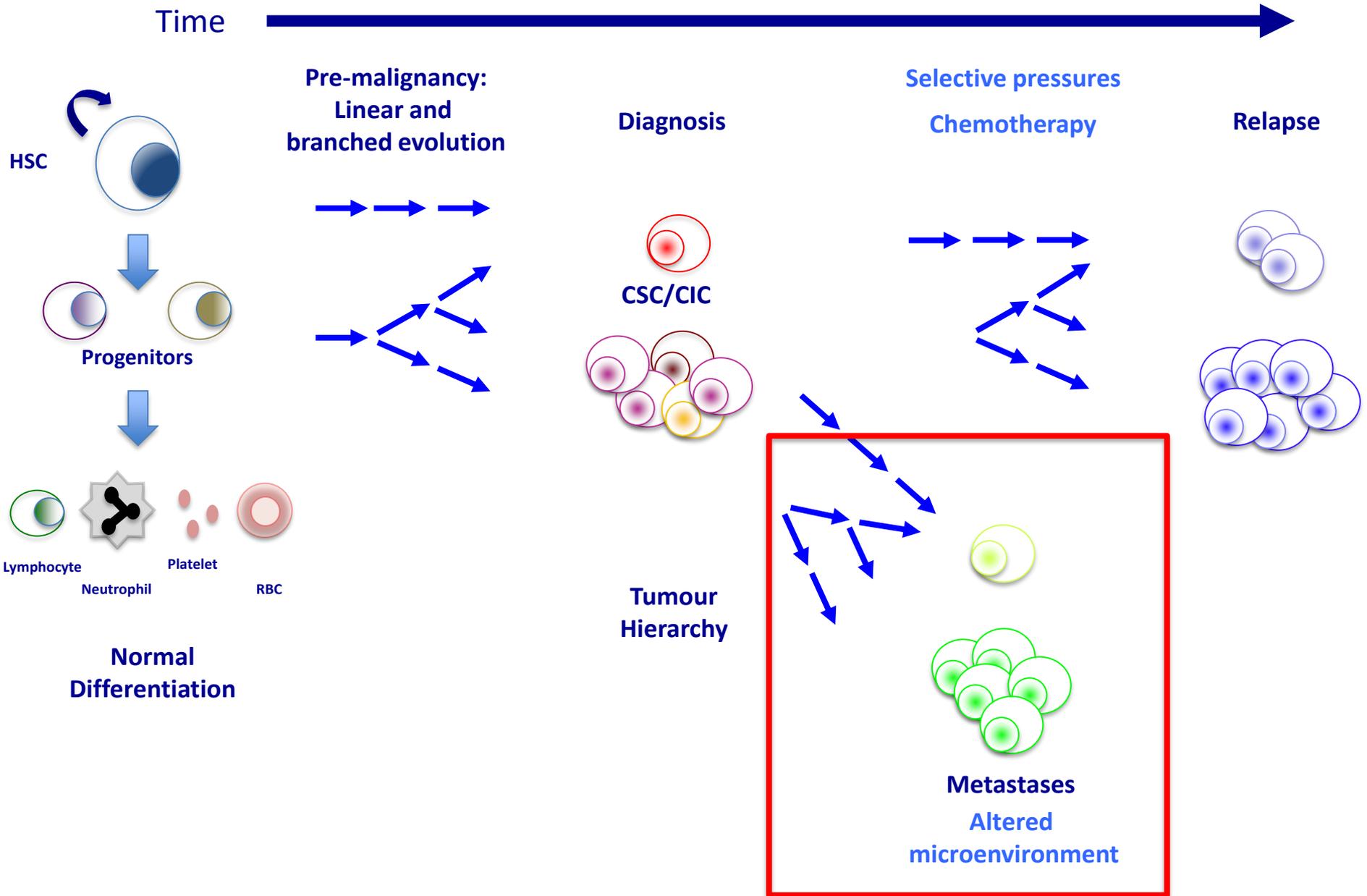
# Patterns of relapse disease also demonstrates temporal intratumoural heterogeneity

## B. Heterogeneous clonal pattern of relapse in AML



- Pattern of relapse reflects significant intratumoural heterogeneity at diagnosis
- Provides further temporal heterogeneity
- Multiple patterns of relapse – major clone, minor clone, evolution of an ancestral clone

# Heterogeneity in Cancer- definition of terms

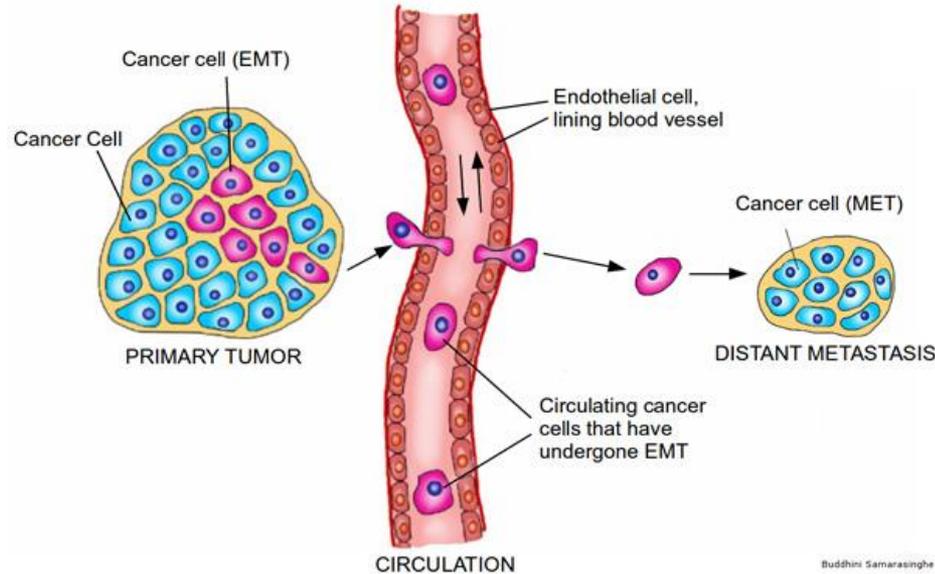


# Fatal consequences of cancer: Metastases and relapse

- We are actually pretty successful at treating primary cancers
- Surgery
- Radiotherapy
- Chemotherapy
- Immunotherapy
- NOT at preventing spread – Metastasis
- NOT at preventing recurrence – Relapse
- Most patients die from relapse and/or spread of disease

# Fatal consequences of cancer: Metastasis

- Metastasis – process of spread of cancer



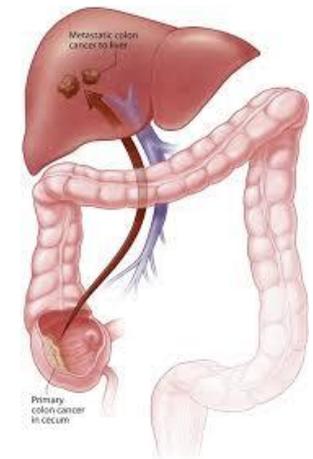
Metastases have differing genetics and functional properties to primary tumours

NEED CSC FUNCTION

MAY DEVELOP HETEROGENEITY FROM PRIMARY TUMOUR  
DUE TO DIFFERENT SELECTIVE PRESSURES OF MICROENVIRONMENT



Seed and soil – different  
Tumours have varying  
predilections for spread

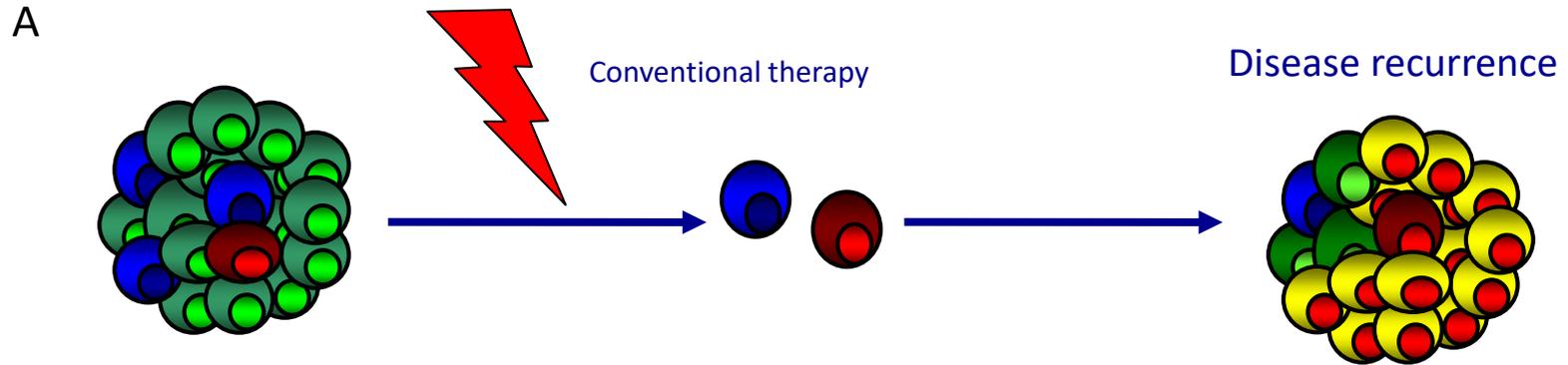


# Therapeutic implications

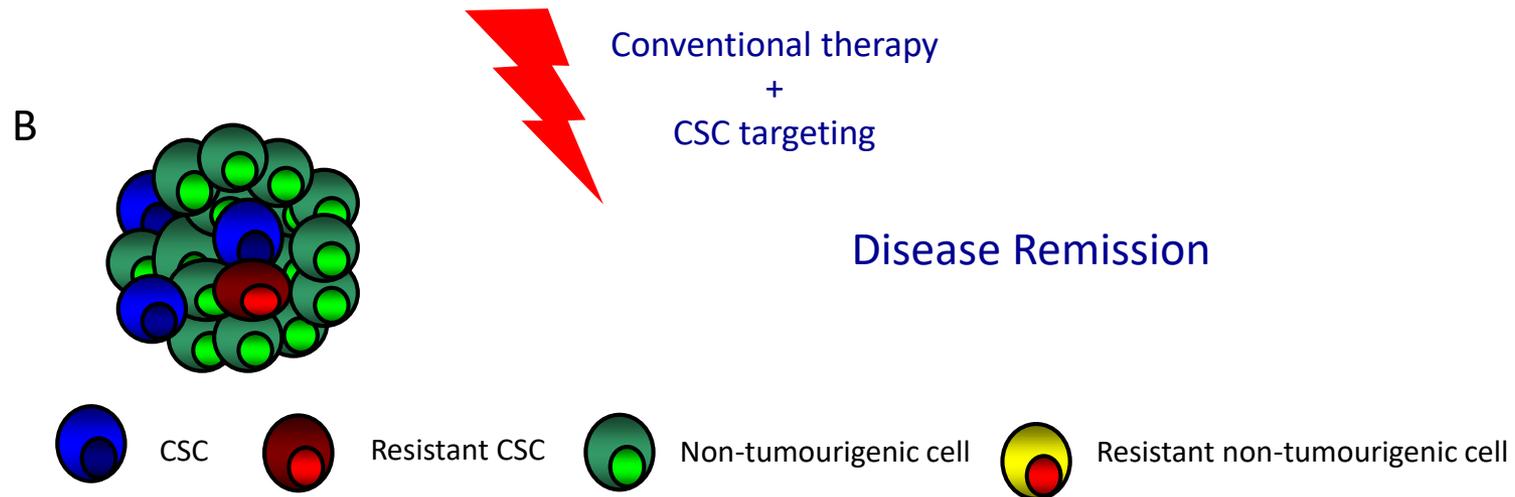
- Targeting cancer stem cells
- Targeting tumour heterogeneity

# TARGETING CSC

## Clinical implications of Cancer Stem Cell model



Targeted elimination of Cancer stem cells (CSC)



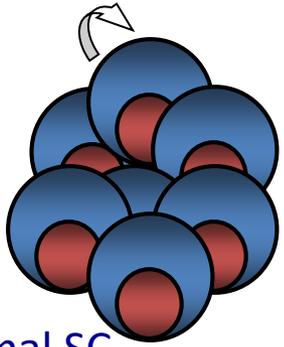
SIMILAR FOR TARGETING HETEROGENEITY

# Targeting cancer stem cells

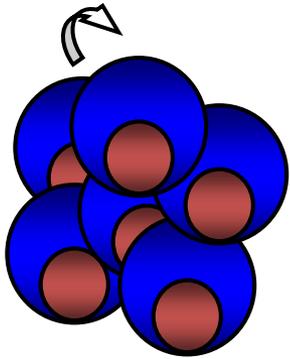
## Considerations:

- Selectivity/Specificity for CSC
- Limited toxicity against normal stem cells from same (or other) organs
- Darwinian selection pressures inherent in CSCs– “MOVING TARGET”
- Need to have an extensive knowledge of the CSC AND its normal counterpart
- Signalling
- Transcriptional/epigenetic
- Metabolic
- Surface proteome
- Immunogenetics

# Identifying differences between normal and cancer stem cells

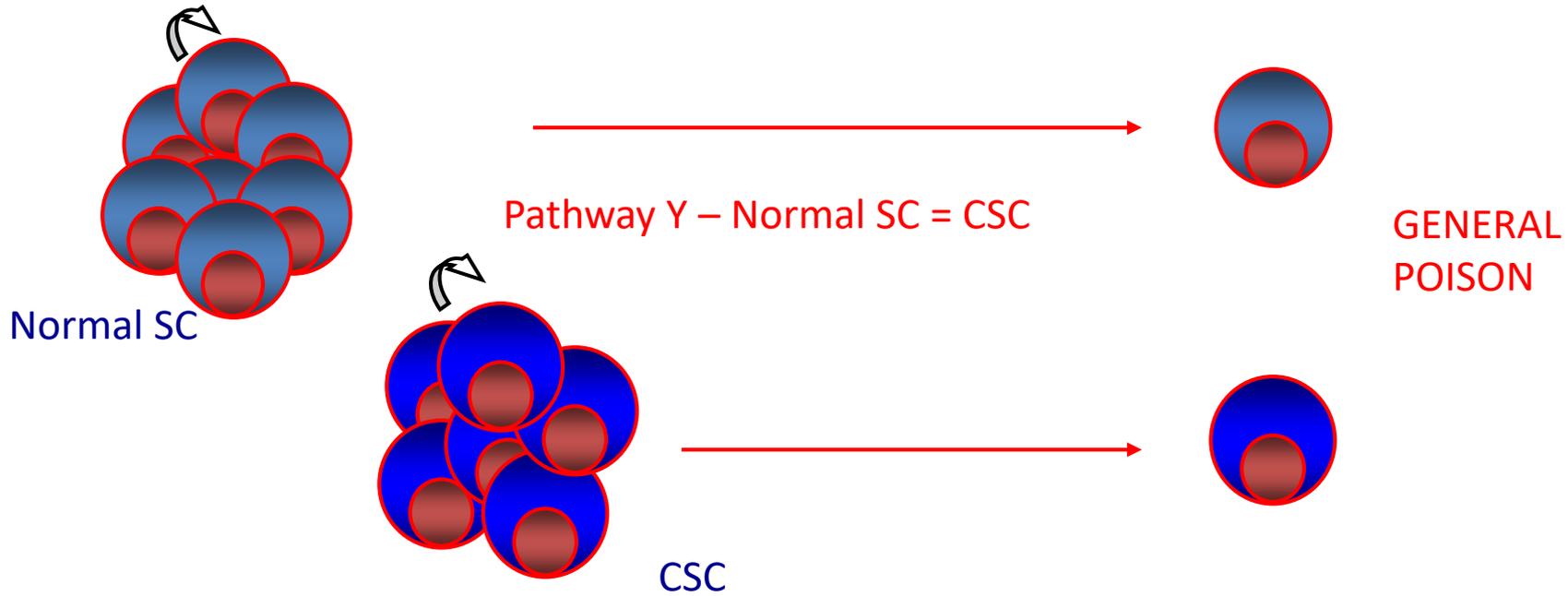


Normal SC

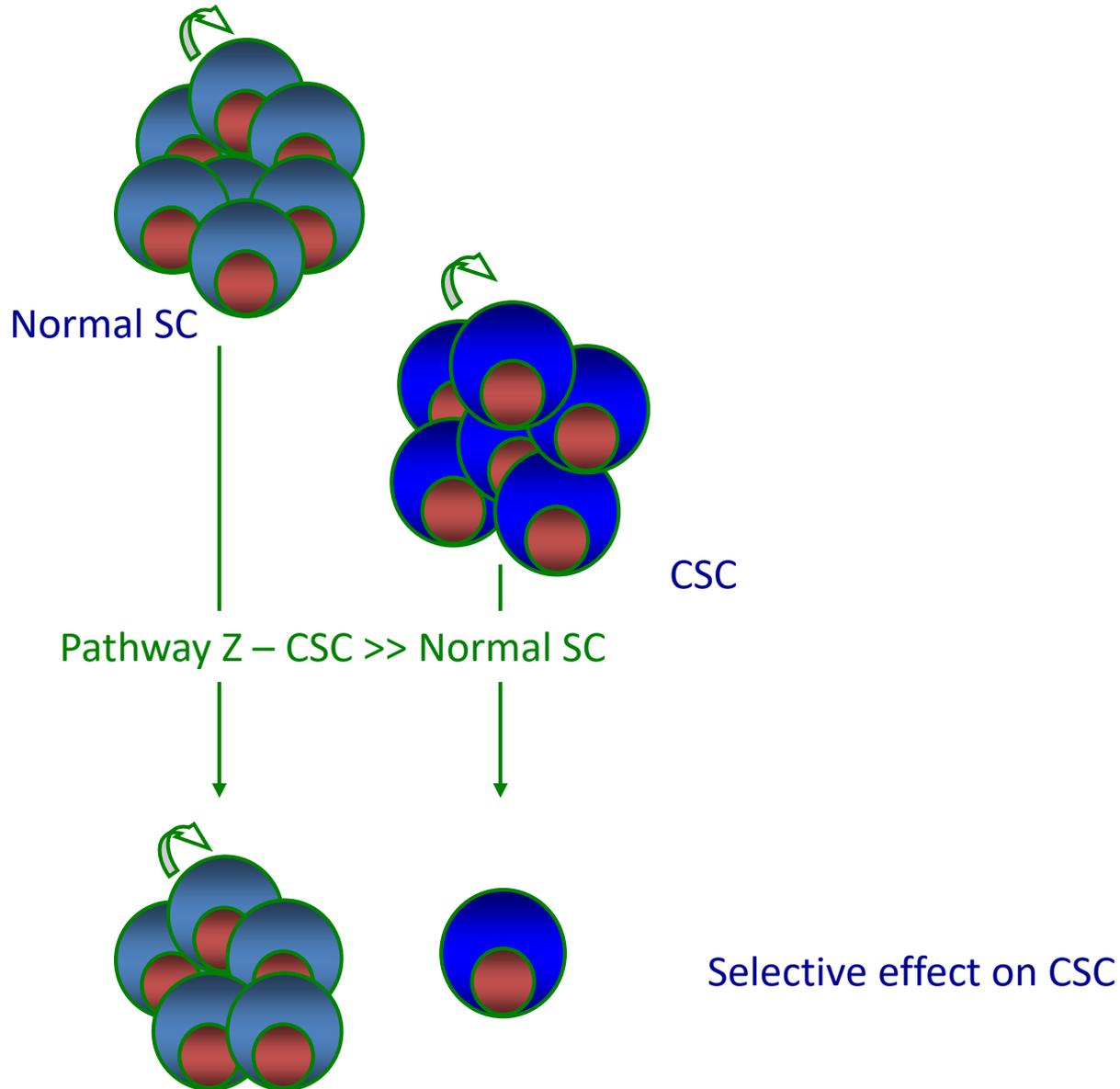


CSC

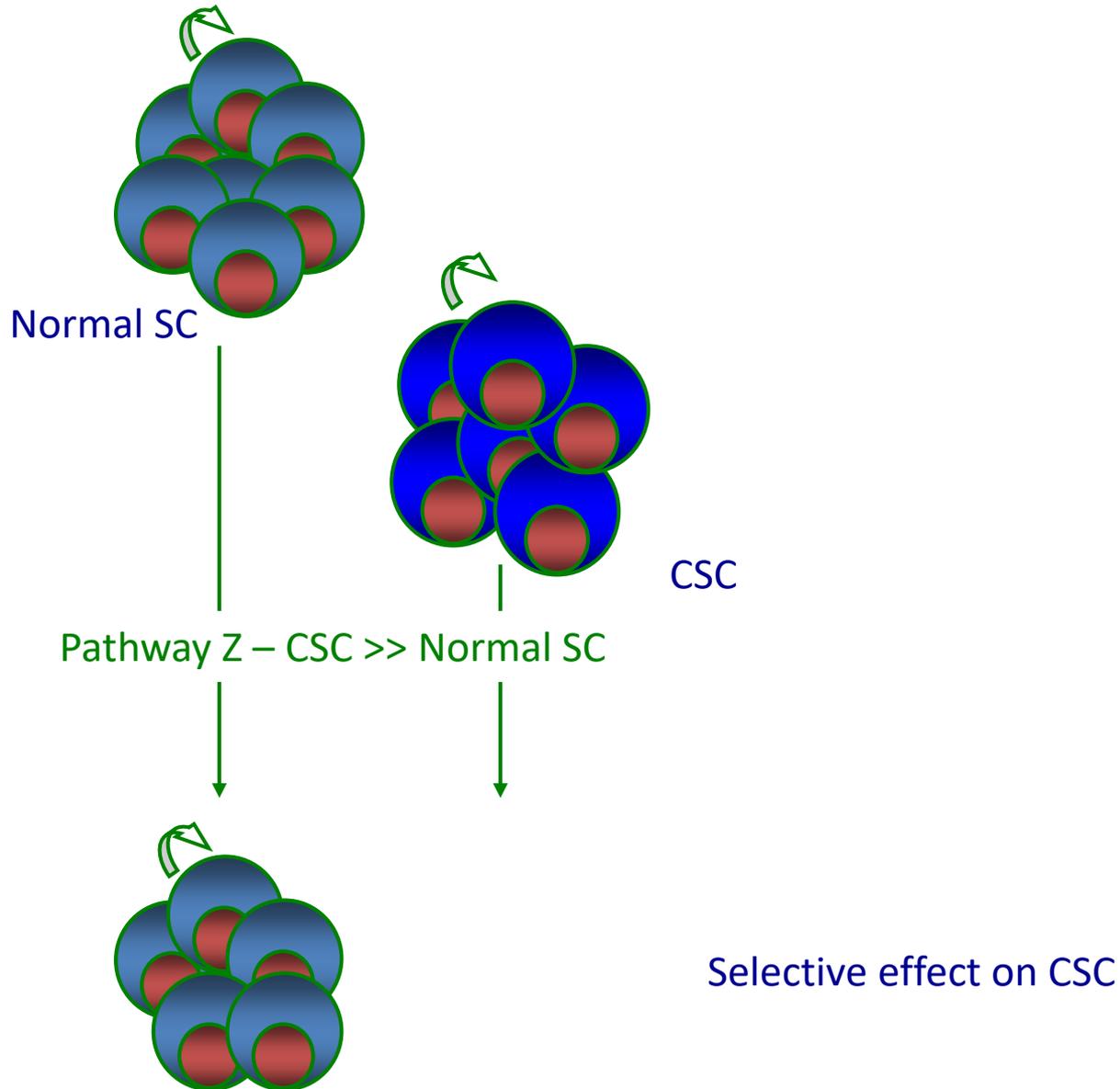
# Identifying differences between normal and leukemic stem cells



# Identifying differences between normal and leukemic stem cells



# Identifying differences between normal and leukemic stem cells



# Targeting LSC surface phenotype: CD47



**CD47 Is Upregulated on Circulating Hematopoietic Stem Cells and Leukemia Cells to Avoid Phagocytosis**

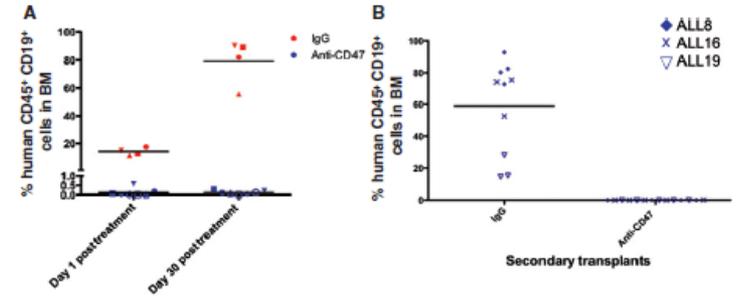
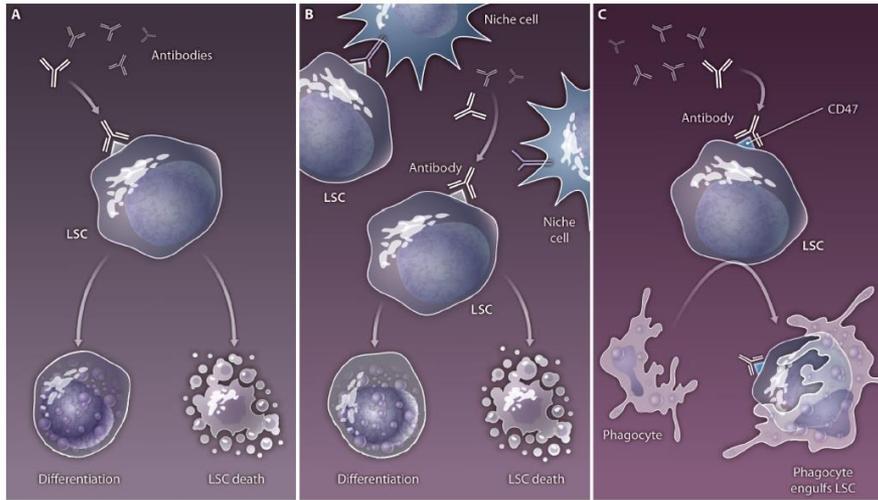
**CD47 Is an Adverse Prognostic Factor and Therapeutic Antibody Target on Human Acute Myeloid Leukemia Stem Cells**

Cell 2009

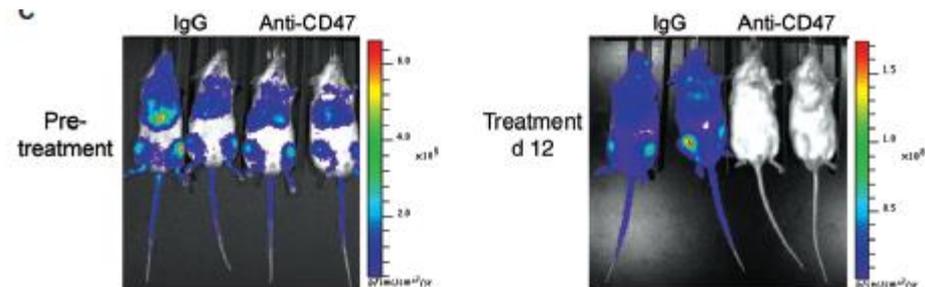
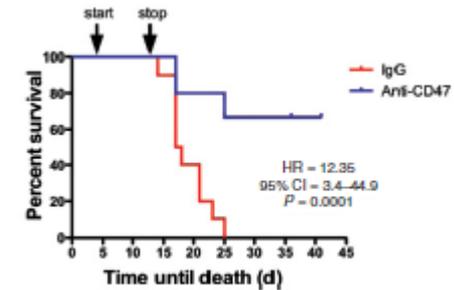
<b>Normal hematopoietic stem cell</b>				
	Setting	Homeostatic maintenance	Inflammation/mobilization	HSC senescence
	CD47 expression	Physiological	Increased	Reduced/absent
	Macrophage activity	Physiological	Increased	Physiological
Net result	No phagocytosis	No phagocytosis	Increased phagocytosis	
Experimental transplantability	Normal	High	Low	
<b>Leukemic stem cell</b>				
	Setting	AML with good prognosis	AML with poor prognosis	AML with poor prognosis treated with anti-CD47
	CD47 expression	Low	High	High (but blocked)
	Macrophage activity	Physiological	Suppressed	Physiological or increased
Net result	Enhanced phagocytosis	No phagocytosis	Enhanced phagocytosis	
Experimental transplantability	Low	Very high	Low	
Clinical prognosis	Good	Very poor	Improved	

- CD47 a “don’t eat me” signal
- Binds to SIRP $\alpha$  on Macrophages and prevents phagocytosis
- During physiology –protective to circulating HSC
- Upregulated in AML vs HSC – selectively protective to LSC
- Potential target

# Targeting LSC surface phenotype: Anti-CD47 preclinical data in ALL



- Preclinical effects vs Acute Lymphoblastic leukaemia (ALL)
- Decreased human engraftment (flow)
- Decreased human engraftment (IVIS- bioluminescence)
- Increased survival



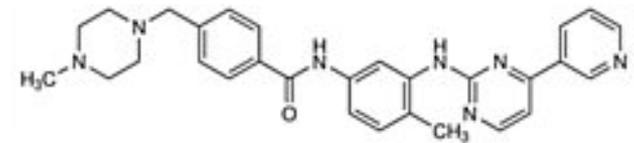
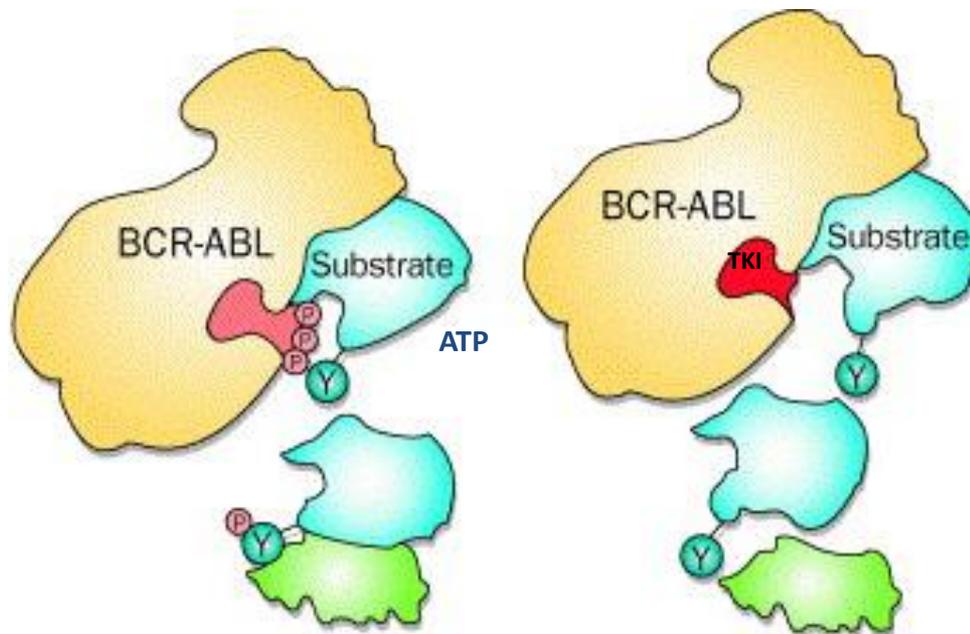
# Therapeutic implications

- Targeting cancer stem cells
- Targeting tumour heterogeneity

# Targeted therapy: Glivec for Chronic Myeloid Leukaemia (CML)

- BCR-ABL gene product, from t (9;22) chromosomal translocation – causal in ALL CASES of CML – little genetic heterogeneity
- Constitutively active Tyrosine kinase
- Competitive inhibition of ATP-binding pocket possible

Prototype compound – Imatinib Mesylate

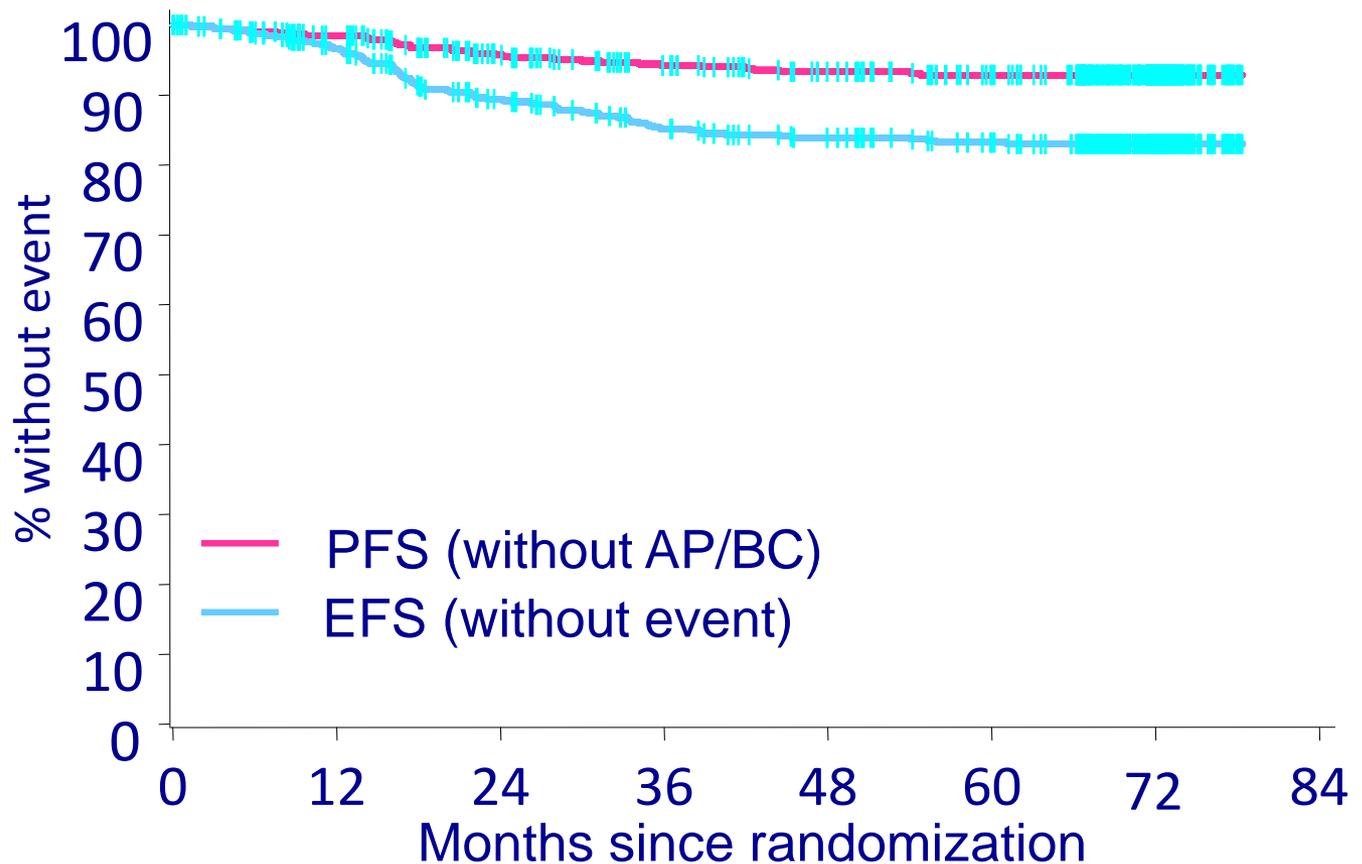


2-phenylaminopyridine derivative

# Targeted therapy: Gleevec for Chronic Myeloid Leukaemia (CML)



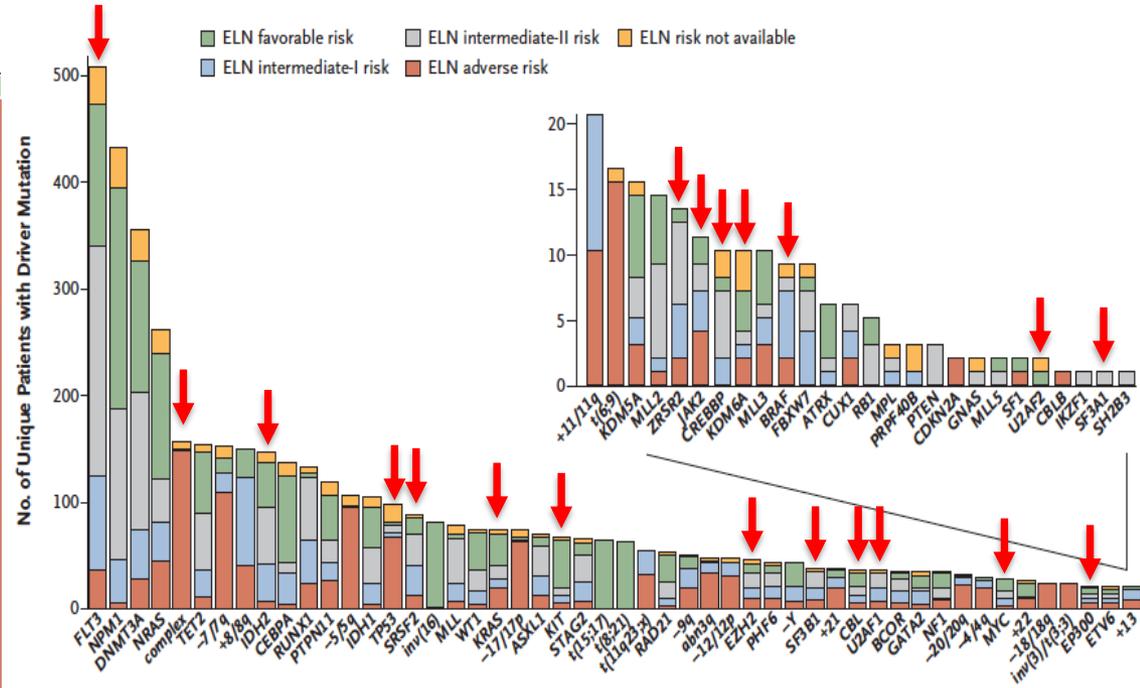
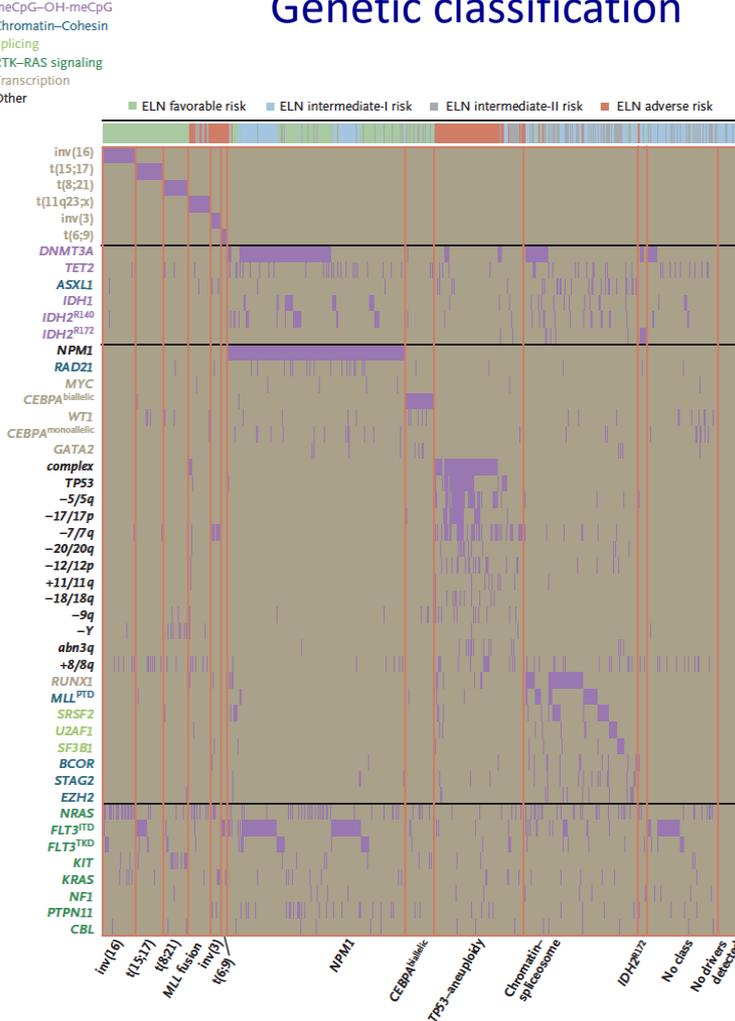
- Previous median survival for CML ~4-5 years
- Was the commonest indication for BMT



WORKS SO WELL DUE TO HOMOGENEITY OF CML –SINGLE MOLECULAR ABNORMALITY

# Heterogeneity between tumours: the basis of personalised medicine

## Genetic classification



Multiple intuitive targets identified from mutational analysis:

- *FLT3*
- *IDH2*
- *SRSF2/U2AF1/U2AF2*
- *KIT*
- *JAK2*
- *BRAF*

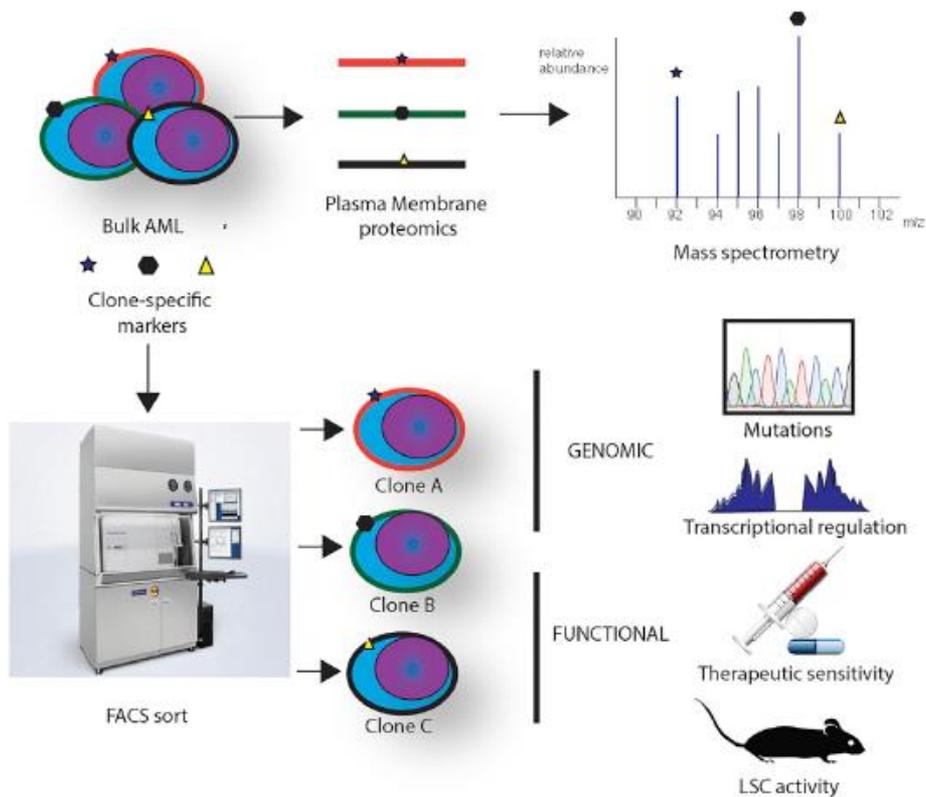
Many mutations SUBCLONAL  
Overt disease  
Occur late during tumour  
devolution

**HETEROGENEITY WITHIN TUMOURS ALSO – TARGET TRUNCAL/INITIATING LESIONS**

# Investigation of intra-tumoural heterogeneity

## Prospective Isolation and Characterization of Genetically and Functionally Distinct AML Subclones

Bauke de Boer,<sup>1</sup> Janine Prick,<sup>1,7</sup> Maurien G. Pruis,<sup>1,7</sup> Peter Keane,<sup>2</sup> Maria Rosaria Imperato,<sup>2</sup> Jennifer Jaques,<sup>1</sup> Annet Z. Brouwers-Vos,<sup>1</sup> Shanna M. Hogeling,<sup>1</sup> Carolien M. Woolthuis,<sup>1</sup> Marije T. Nijk,<sup>3</sup> Arjan Diepstra,<sup>4</sup> Sebastian Wandinger,<sup>5</sup> Matthias Versele,<sup>6</sup> Ricardo M. Attar,<sup>6</sup> Peter N. Cockerill,<sup>2</sup> Gerwin Huls,<sup>1</sup> Edo Vellenga,<sup>1</sup> André B. Mulder,<sup>3,8</sup> Constanze Bonifer,<sup>2,8</sup> and Jan Jacob Schuringa<sup>1,8,9,\*</sup>



## Cancer Cell Article

- Used plasma membrane proteomics (PMP) to identify AML specific surface proteins
- Some discriminatory for individual clones within specific patients
- Used to isolate clones – Different Biology
- Link to LSC potential (CD25)
- Link to genotype (CD25 and FLT3-ITD)
- Different regulation of transcription
- Different response to drugs

**Intratumoral Heterogeneity: Tools to Understand and Exploit Clone Wars in AML**

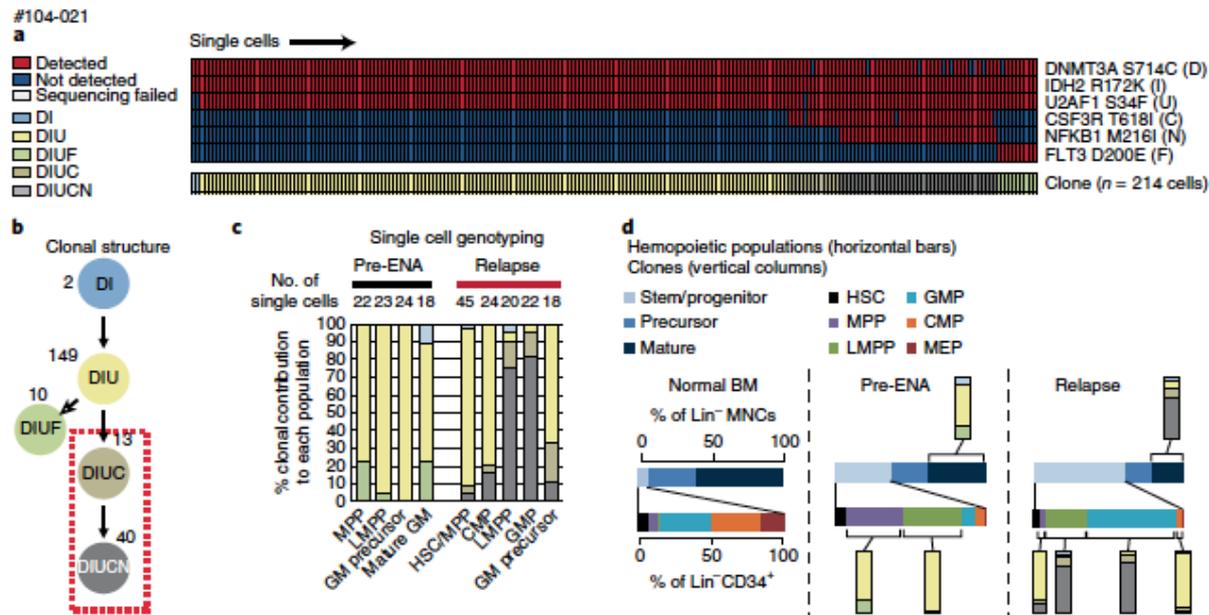
George Giotopoulos<sup>1,2,3</sup> and Brian J.P. Huntly<sup>1,2,3,\*</sup>

# Investigation of intra-tumoural heterogeneity: Relapse

## Clonal heterogeneity of acute myeloid leukemia treated with the IDH2 inhibitor enasidenib



Lynn Quek<sup>1,2,3,16\*</sup>, Muriel D. David<sup>4,16</sup>, Alison Kennedy<sup>1,2</sup>, Marlen Metzner<sup>1,2</sup>, Michael Amatangelo<sup>5</sup>, Alan Shih<sup>6</sup>, Bilyana Stoilova<sup>1,2</sup>, Cyril Quivoron<sup>4</sup>, Maël Heiblig<sup>4</sup>, Christophe Willekens<sup>4,7</sup>, Véronique Saada<sup>4,7</sup>, Samar Alsafadi<sup>8</sup>, M. S. Vijayabaskar<sup>9</sup>, Andy Peniket<sup>3</sup>, Oliver A. Bernard<sup>4</sup>, Sam Agresta<sup>10</sup>, Katharine Yen<sup>10</sup>, Kyle MacBeth<sup>5</sup>, Eytan Stein<sup>6</sup>, George S. Vassiliou<sup>9,11,12,13</sup>, Ross Levine<sup>6,14,15</sup>, Stephane De Botton<sup>4,7,17\*</sup>, Anjan Thakurta<sup>5,17\*</sup>, Virginie Penard-Lacronique<sup>4,17\*</sup> and Paresh Vyas<sup>1,2,3,17\*</sup>



- Single cell clonal analysis to determine genetic mechanisms of relapse
- Clonal evolution
- Selection of ancestral or terminal clones

# Summary of salient points

- Intra-tumoural vs Inter-tumoural heterogeneity
- Sources of heterogeneity – cell of origin, multiple selective pressures
- Phases during cancer evolution – premalignancy, diagnostic disease, relapsed disease, metastases
- Constant Darwinian nature of tumour evolution – MOVING TARGET
- To target CSC we need to understand them better
- Need to target early/initiating mutations
- A better understanding of early tumour development should allow for earlier (and more successful) therapeutic intervention