

Methods in Haematopoietic Stem Cell Biology

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MRC WIMM DPhil Course November 2019

Learning Objectives

- What is a **haematopoietic stem cell (HSC)**?
- **Key concepts** in HSC biology during ontogeny and adult life
 - Definitions and properties of HSCs and progenitors
 - HSC niches
 - Intrinsic / extrinsic regulators
- **Experimental approaches** to study HSC biology
 - *In vitro*
 - *In vivo*
- **Cancer Stem cells**

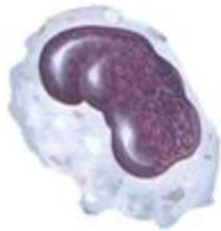
What is haematopoiesis?

The process that generates blood cells
of all lineages throughout life
(from 3 weeks gestation)

Blood cell lineages

Blood Cells

White cells



Monocyte



Lymphocyte



Neutrophil



Eosinophil



Basophil

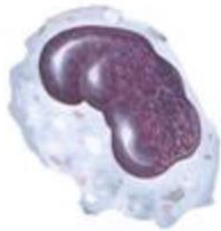


Macrophage

Blood cell lineages

Blood Cells

White cells



Monocyte



Lymphocyte



Neutrophil



Eosinophil



Basophil



Macrophage



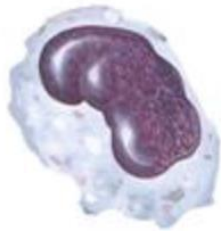
Erythrocyte

Red cells

Blood cell lineages

Blood Cells

White cells



Monocyte



Lymphocyte



Neutrophil



Eosinophil



Basophil



Macrophage



Erythrocyte



Platelets

Platelets

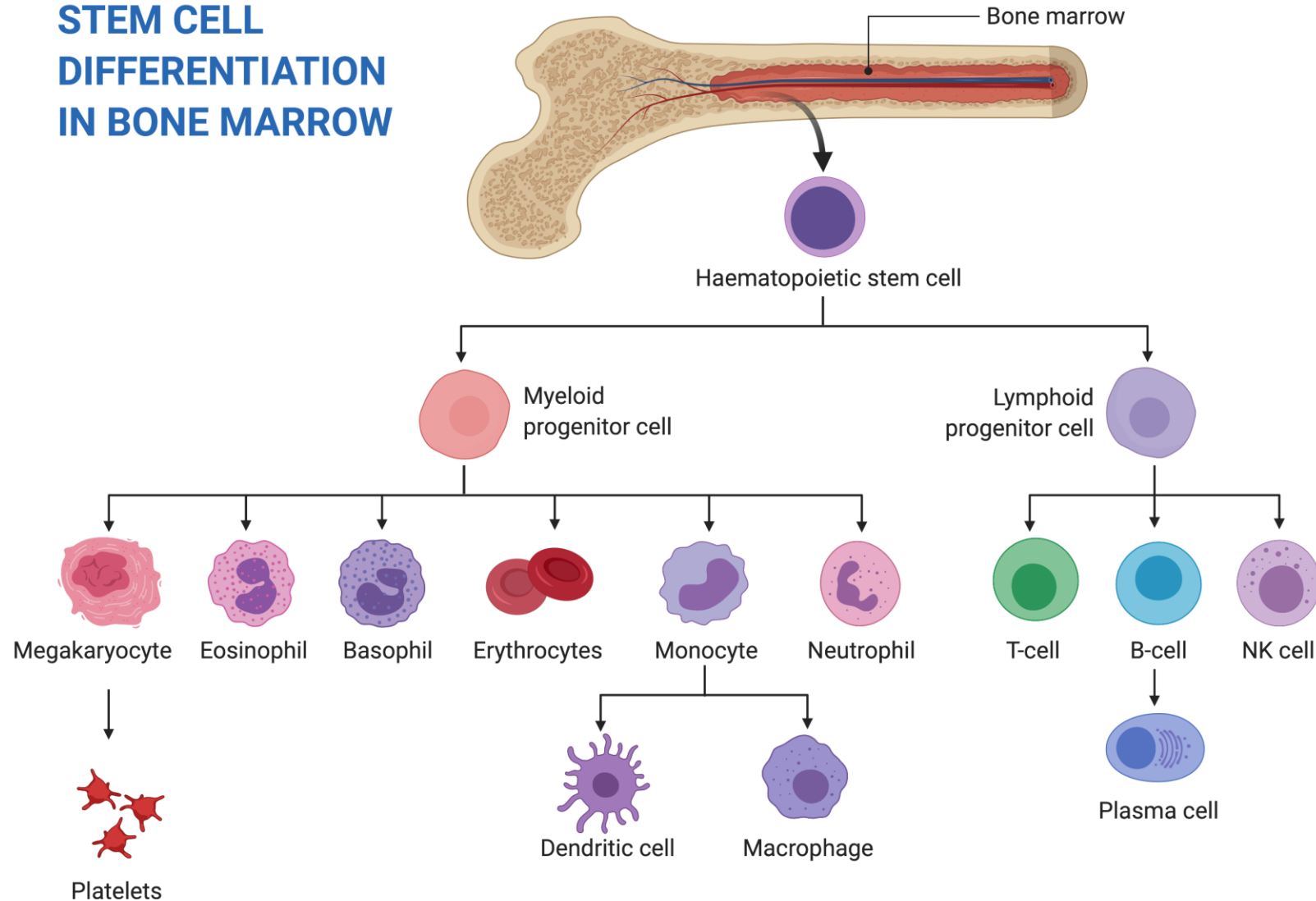
Red cells

HSCs are arguably the best studied stem cell in mammals

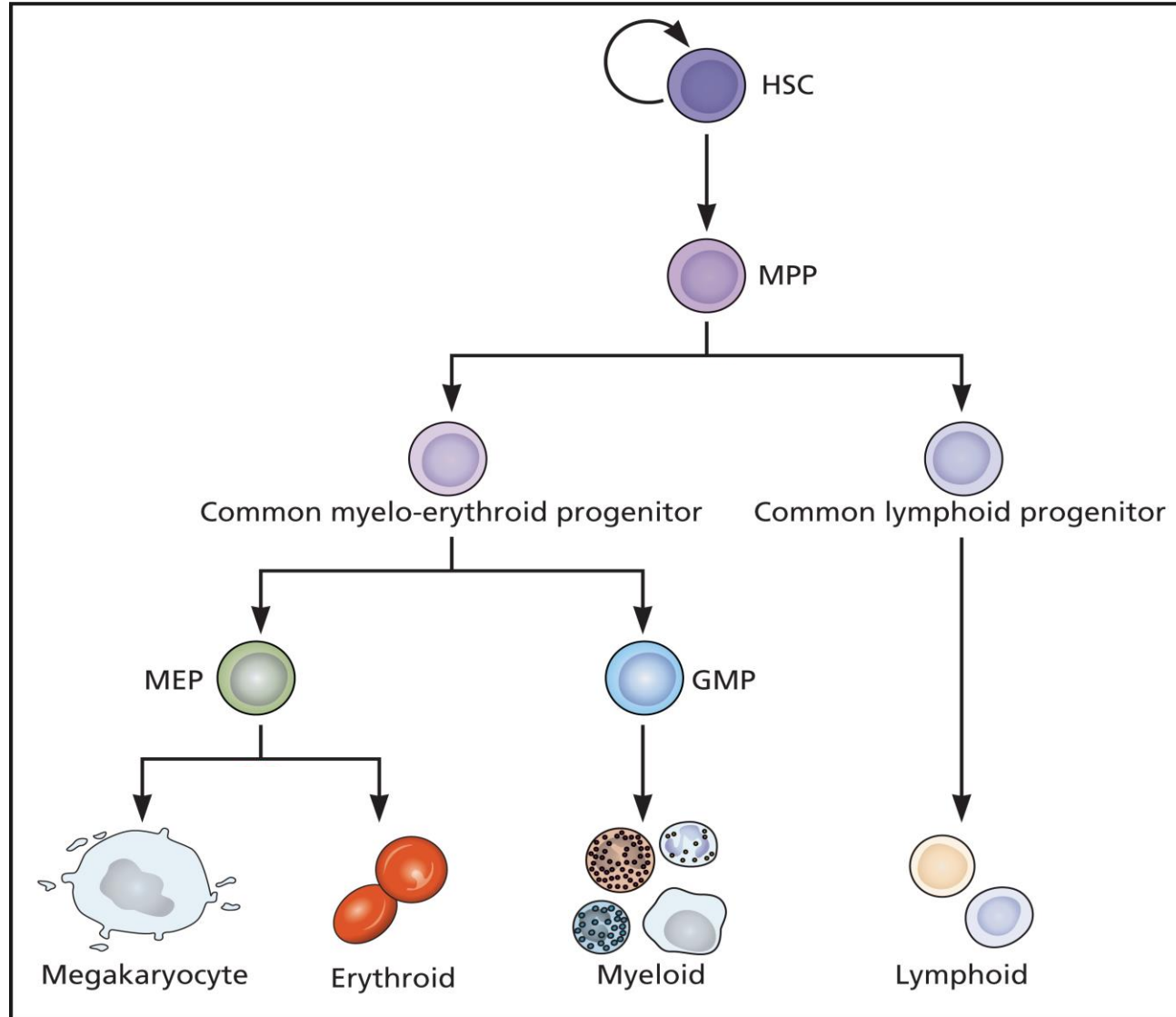
- Easy of access
- Blood lineages are morphologically and immunophenotypically distinct
- Experimental assays now well developed
- Only stem cell therapy in routine clinical use
- While stem cell transplant is widely used, there are many limitations
 - Mortality <25%
 - Graft-versus-host-disease
 - Relapse

Haematopoiesis: classical hierarchical model

STEM CELL DIFFERENTIATION IN BONE MARROW

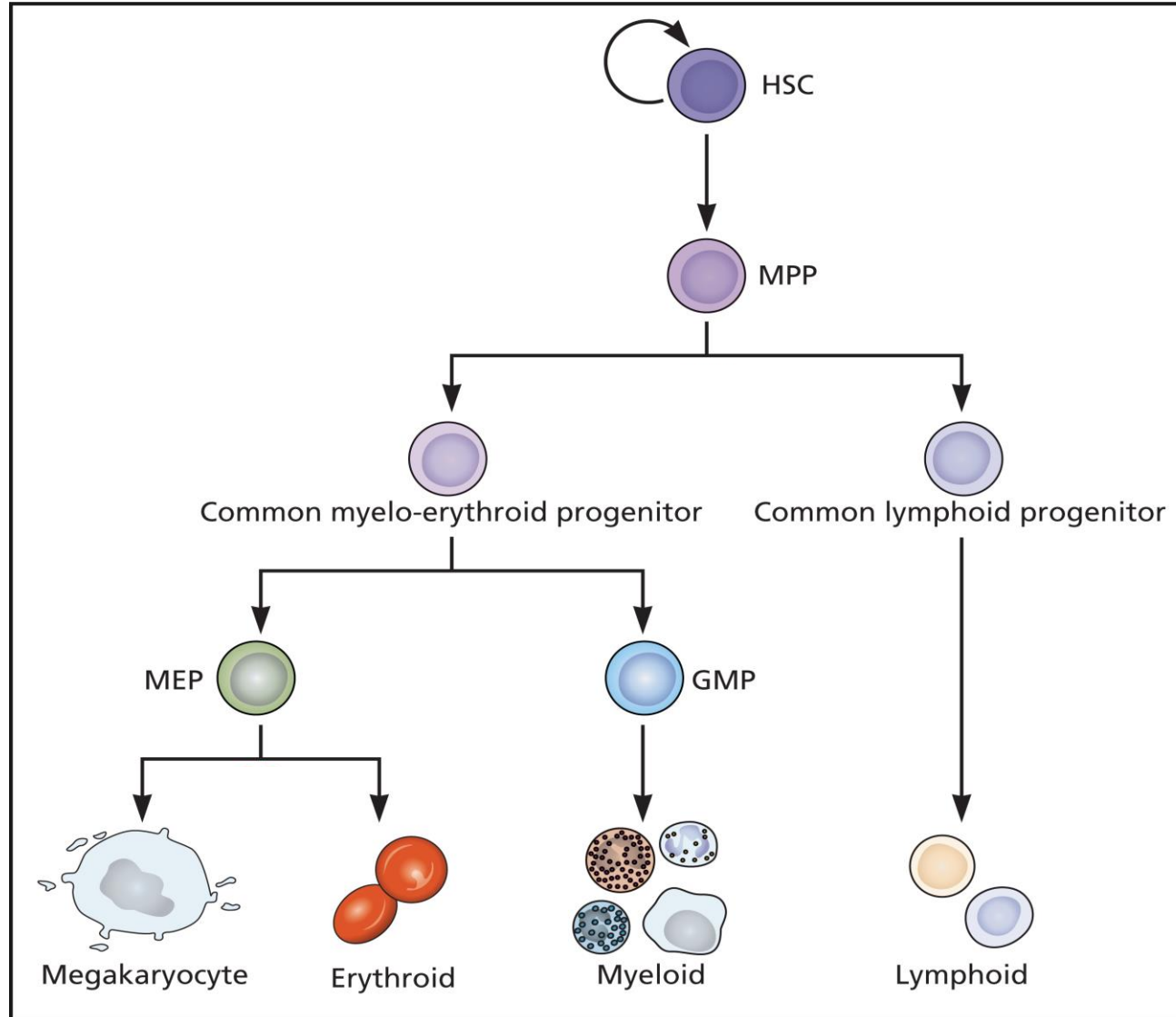


Haematopoietic stem cells



What are the key defining properties of an HSC?

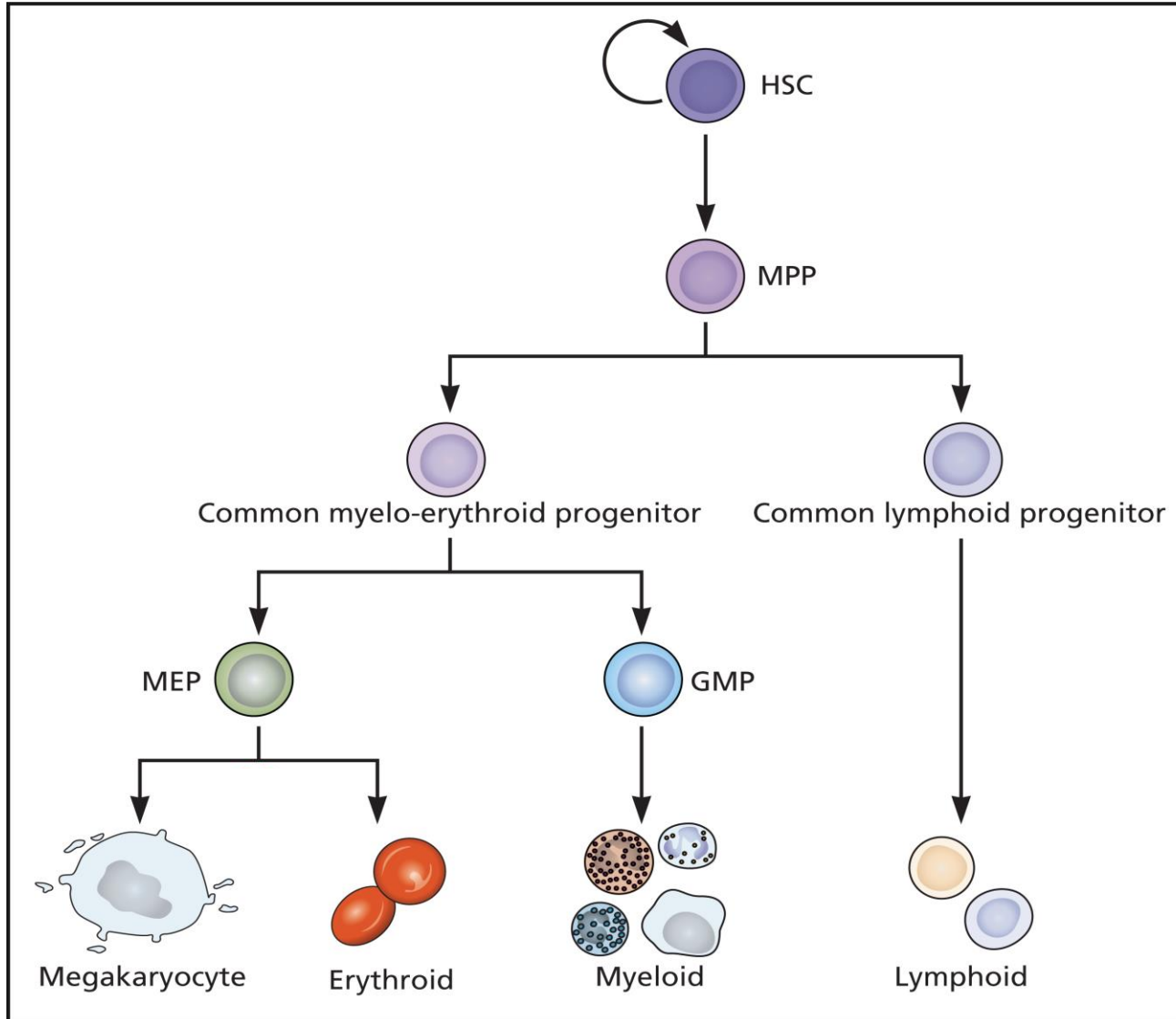
Haematopoietic stem cells



Properties of HSCs

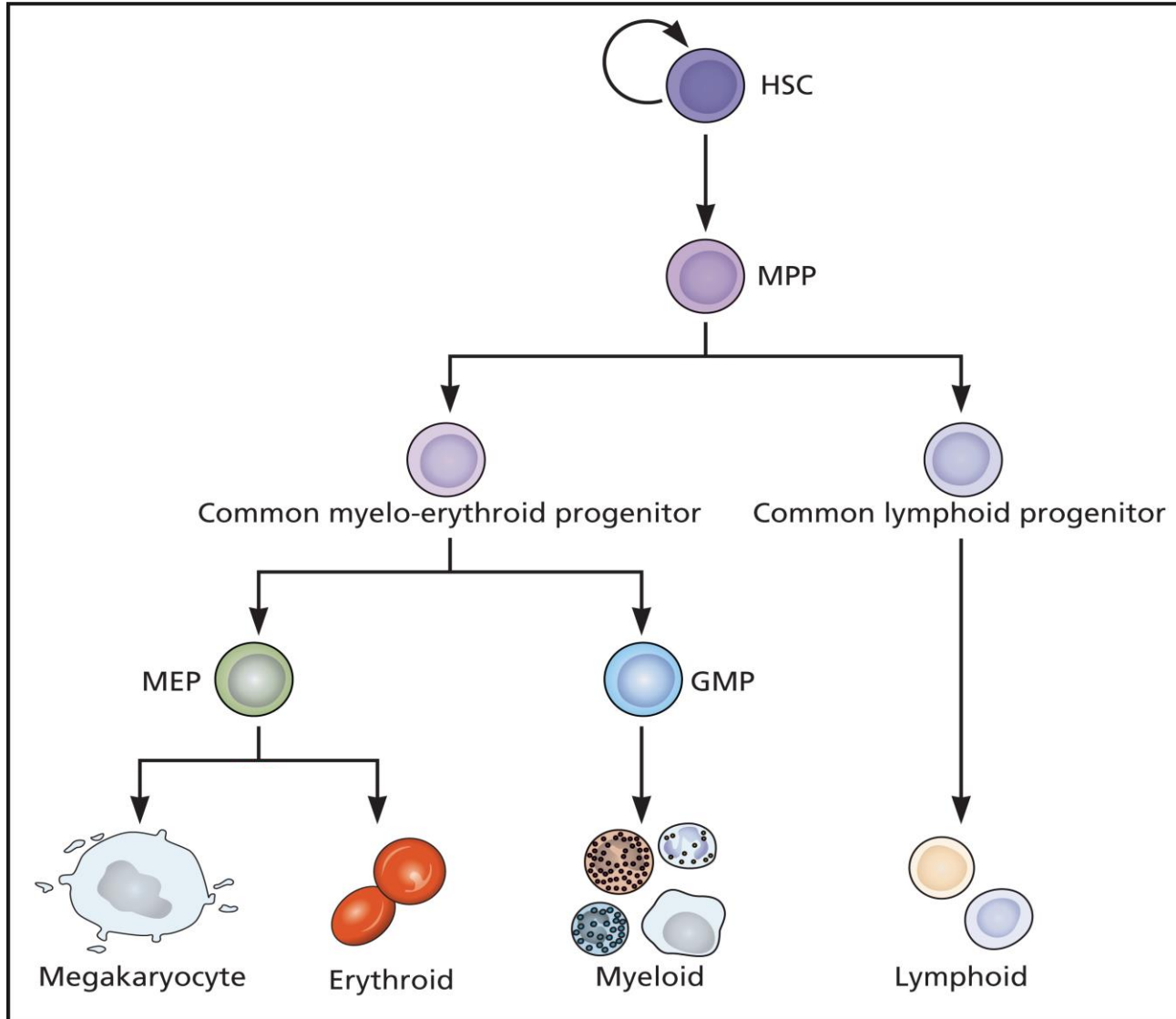
- Self renewal
- Multi-lineage differentiation
- When transplanted, can reconstitute life-long haematopoiesis

Haematopoietic progenitor cells



- Less self-renewal capacity (short term engraftment)
- Already made some lineage commitment decisions

Haematopoietic progenitor cells



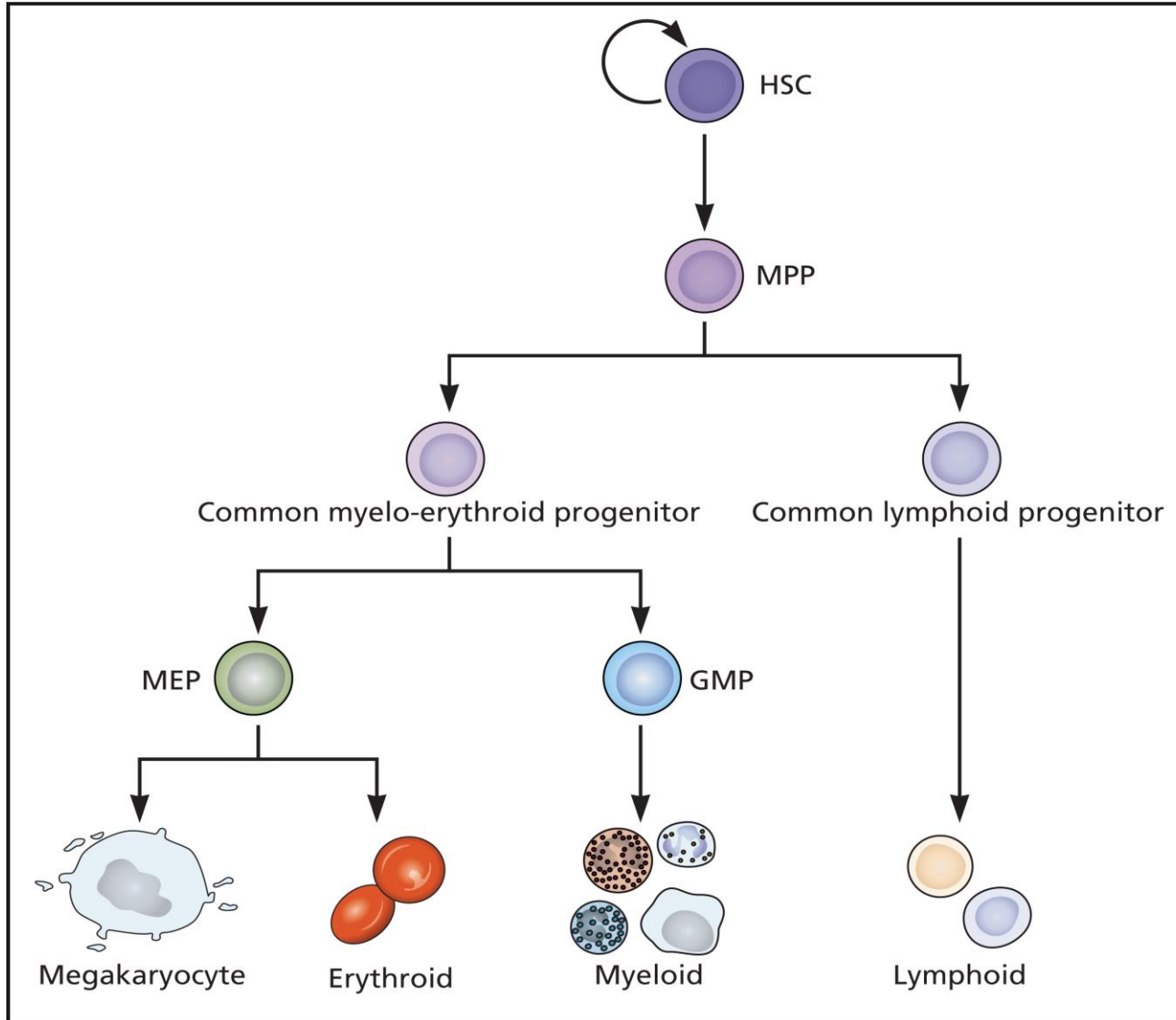
- Common lymphoid progenitor

(B, T, NK cells)

- Common myeloid progenitor

(granulocytes, erythroid, megakaryocyte and monocyte/macrophage)

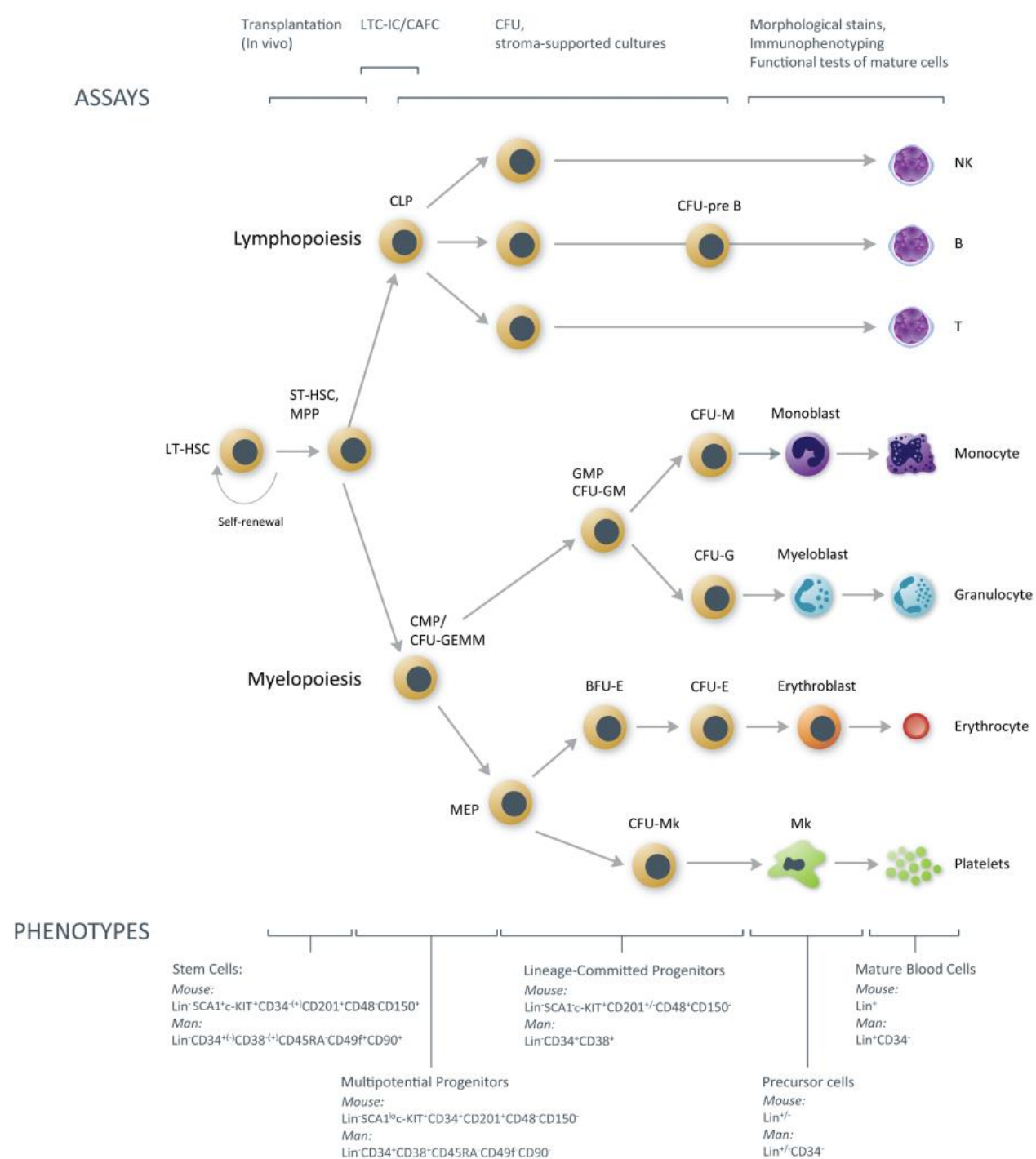
Haematopoietic stem/progenitor cells



**Identified by their
expression of cell surface
markers**

e.g. Human HSC =
CD34+lineage-CD38-
CD45RA-CD90+CD49f+

Identification of HSC/HSPCs: Human/mouse



B

Cell type	Mouse marker		Human marker
HSCs	LT-HSC	Lin ⁻ Sca-1 ⁺ c-Kit ⁺ CD34 ⁻ Flk2 ⁻	CD34 ⁺ CD38 ⁻ CD90 ⁺ CD45RA ⁻ CD49F ⁺
	ST-HSC	Line ⁻ Sca-1 ⁺ c-Kit ⁺ CD34 ⁺ Flt2 ⁻	
	MPPs	Lin ⁻ Sca-1 ⁺ c-Kit ⁺ CD34 ⁺ Flt2 ⁺	CD34 ⁺ CD38 ⁻ CD90 ⁻ CD45RA ⁻ CD49F ⁻
SLAM-HSCs	SLAM-HSC	Lin ⁻ Sca-1 ⁺ c-Kit ⁺ CD150 ⁺ CD48 ⁻ CD229 ⁻ CD244 ⁻	
	SLAM-MPP	Lin ⁻ Sca-1 ⁺ c-Kit ⁺ CD150 ⁻ CD48 ⁻ CD229 ⁻ CD244 ⁻	
Progenitor cells	CLP	Lin ⁻ Flt2 ⁺ IL7Ra ⁺ CD27 ⁺	CD34 ⁺ CD10 ⁺ CD7 ⁺
	CMP	Line ⁻ Sca-1 ⁻ c-Kit ⁺ CD34 ⁺ FcγRII/III ⁻	CD34 ⁺ CD38 ⁺ CD123 ^{med} CD135 ⁺ CD45RA ⁻
	GMP	Lin ⁻ Sca-1 ⁻ c-Kit ⁺ CD34 ⁺ FcγRII/III ⁺	CD34 ⁺ CD38 ⁺ CD123 ^{med} CD135 ⁺ CD45RA ⁺
	MEP	Lin ⁻ Sca-1 ⁻ c-Kit ⁺ CD34 ⁻ FcγRII/III ⁻	CD34 ⁺ CD38 ⁺ CD123 ⁻ CD135 ⁻ CD45RA ⁻ CD110 ⁺
Niche supporting cells	Mφ	CD11b ⁺ Gr1 ^{low} F4/80 ⁺ SSC ^{low}	
	T cell	CD3 ⁺	CD4 ⁺ CD25 ⁺ CD127 ^{low} CD45RA ^{+/-}
	B cell	CD45R/B220 ⁺	
	Erythrocyte	CD45 ⁻ Ter119 ⁺	
	EC	CD45 ⁻ Ter119 ⁻ CD31 ⁺	
	BMSC	CD45 ⁻ Ter119 ⁻ CD31 ⁻ LepR ⁺	CD45 ⁻ CD34 ⁻ CD73 ⁺ CD105 ⁺ CD90 ⁺
	OB	CD45 ⁻ Ter119 ⁻ CD31 ⁻ Sca1 ⁻ CD51 ⁺	

Key markers
HUMAN – CD34, CD38

MOUSE – LSK, CD34-

Table 1

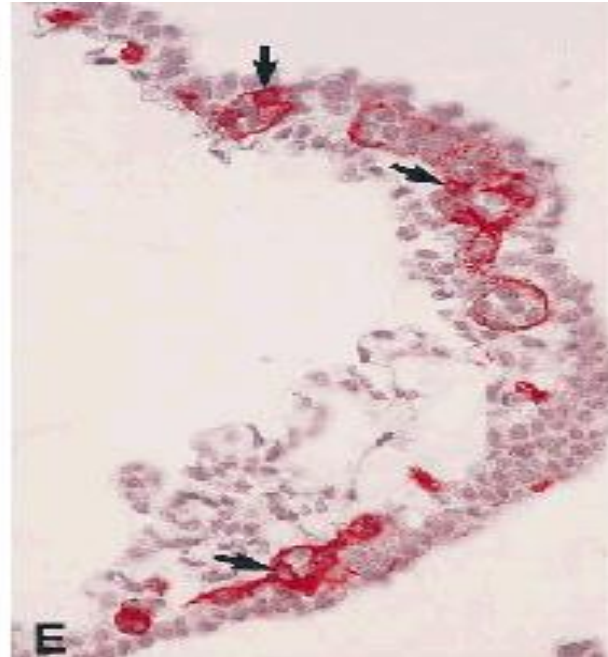
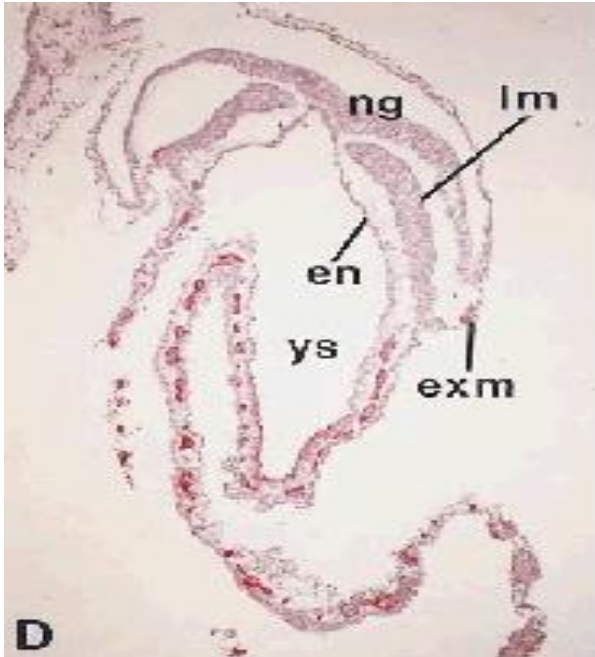
Unique and overlapping commonly used stem cell terms and definitions

Term	Widely accepted definition
LT-HSC	Cells able to reconstitute all five types of mature blood cells in recipient mice over the long term (≥ 16 -weeks after transplantation) and in secondary and tertiary transplants ²⁸
Intermediate term repopulating HSC (IT-HSC)	Cells that reconstitute multi-lineage blood cells over the medium term (6-8 months), but show some loss of self-renewal in secondary transplant recipients ²⁸
Short term repopulating HSC (ST-HSC)	Cells that transiently reconstitute multiple blood cell types for up to 8 weeks in transplant recipients, or in which one or more donor-derived lineages disappear before 24 weeks after primary transplant
Lineage priming	Expression of a transcriptional program associated with potential for differentiation to mature cells of a specific lineage in a multi/oligo-potent cell
Lineage bias	Multi/oligo-potent cells that preferentially give rise to a single lineage but retain potential for alternative differentiation
Lineage restricted/committed	Oligo/unipotent cells not able to give rise to cells of certain lineages
Lineage potential	The mature lineages that a cell <i>has the potential to</i> give rise depending on external stimuli
Lineage fate	The lineage that a stem/progenitor will give rise to <i>in vivo</i>

Sites of haematopoiesis: **embryonic life**

First signs are seen in the **yolk sac** at 3 weeks of gestation

PRIMITIVE HAEMATOPOIESIS

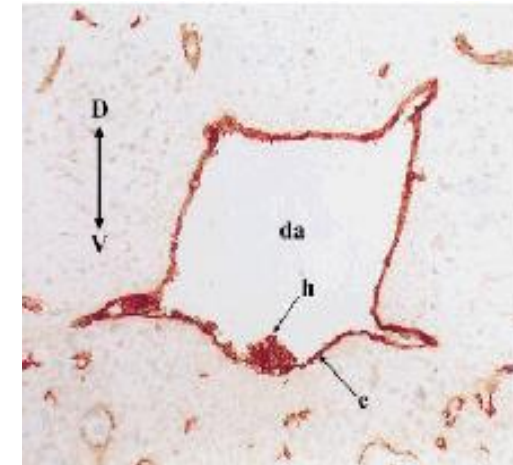
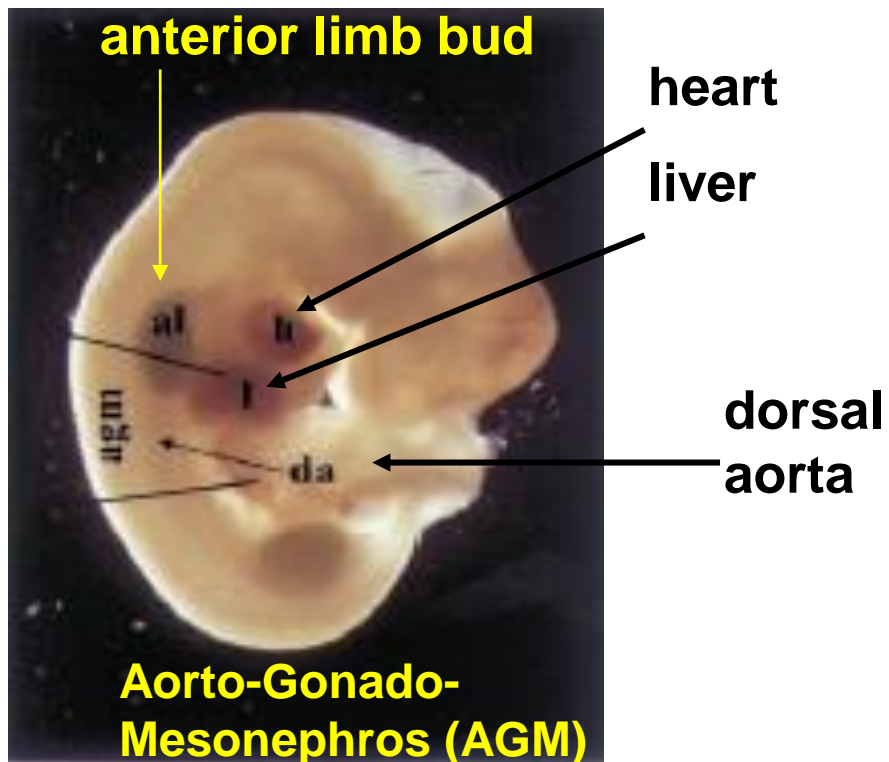


**Immunostaining
with anti-CD34
antibody marks
haemopoietic stem/
progenitors cells &
endothelial cells**

Sites of haematopoiesis: **embryonic life**

DEFINITIVE HAEMATOPOIESIS begins in the aorto-gonado-mesonephros (AGM) at 5 weeks gestation

34-day human embryo

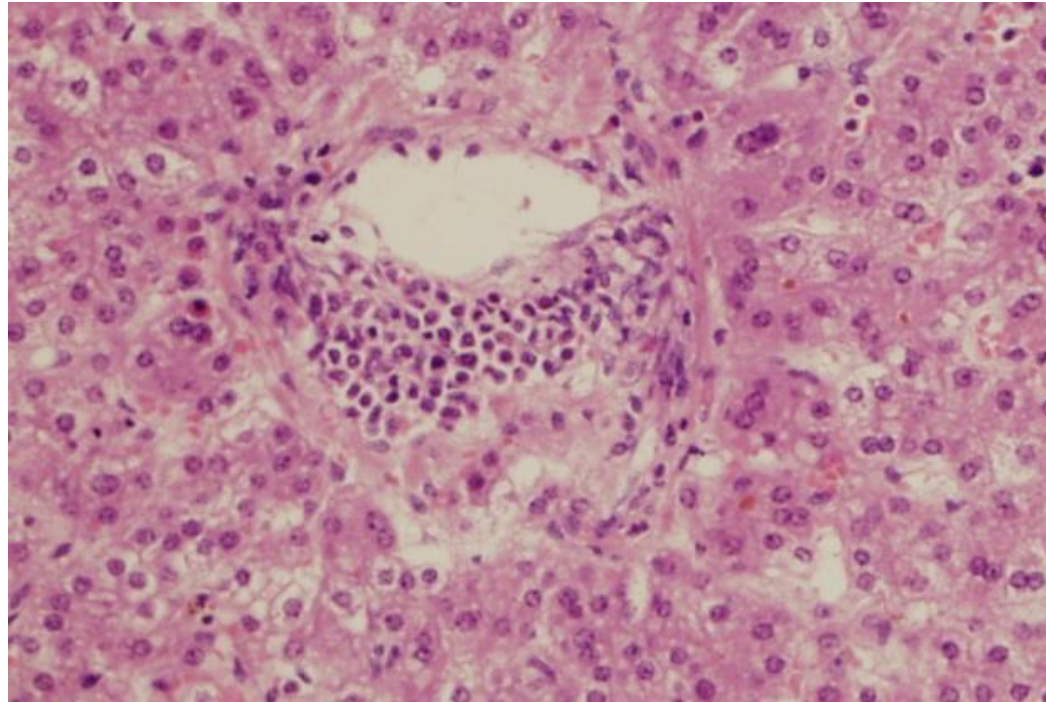


Cross section through the dorsal aorta to show haemopoietic cells (h) in the AGM (stained with CD34)

Sites of haematopoiesis: **embryonic life**

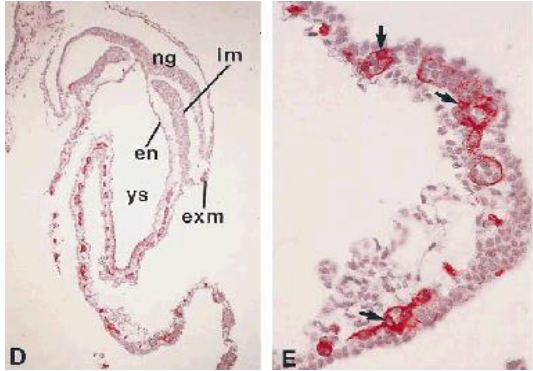
DEFINITIVE HAEMATOTOPOIESIS in the fetal liver begins at 7-8 weeks

This is the main site of haematopoiesis in fetal life



Sites of haematopoiesis: embryonic life

YOLK SAC



Generation of HSC
Mainly erythropoiesis

3 wks

Primitive

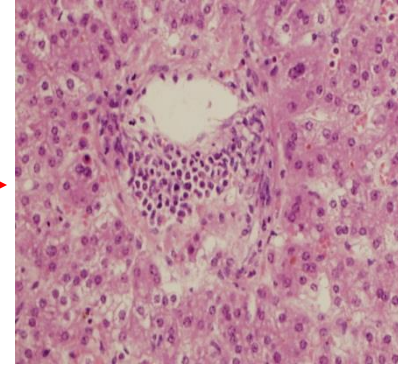
AGM



Generation of HSC
No expansion of HSC

5 wks

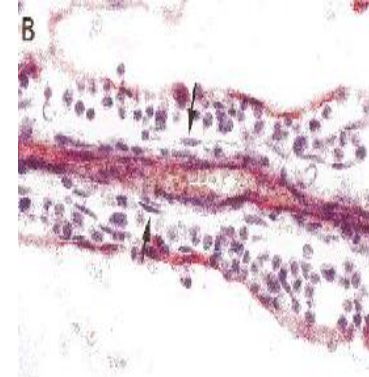
LIVER



HSC maintenance
HSC expansion

7-8 wks

BONE MARROW



Main site from 6-7 months of fetal life and after birth

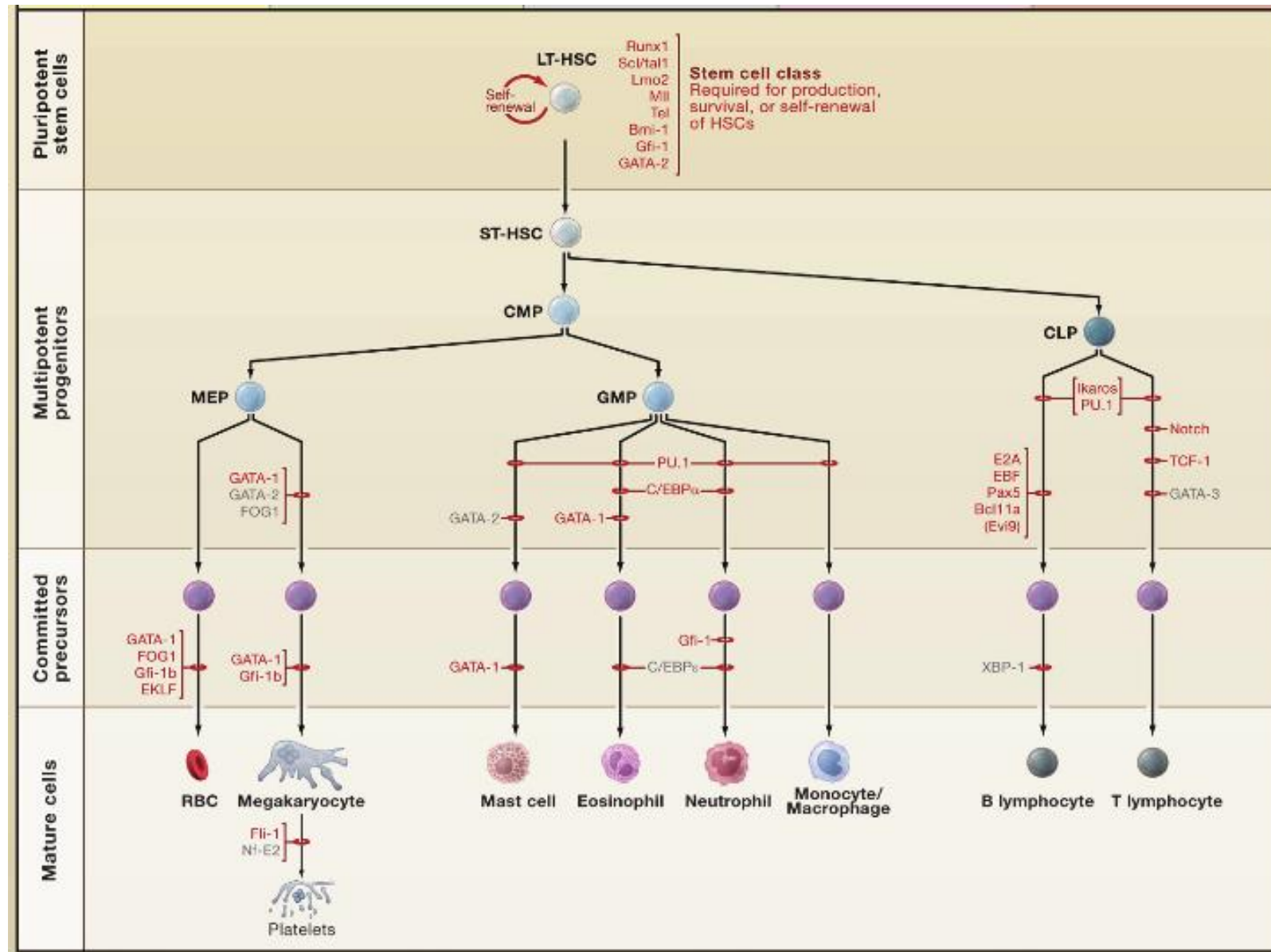
12 wks

Definitive haemopoiesis

Regulation of haematopoiesis

- Cell intrinsic
- Cell extrinsic

Cell intrinsic controls on haematopoiesis



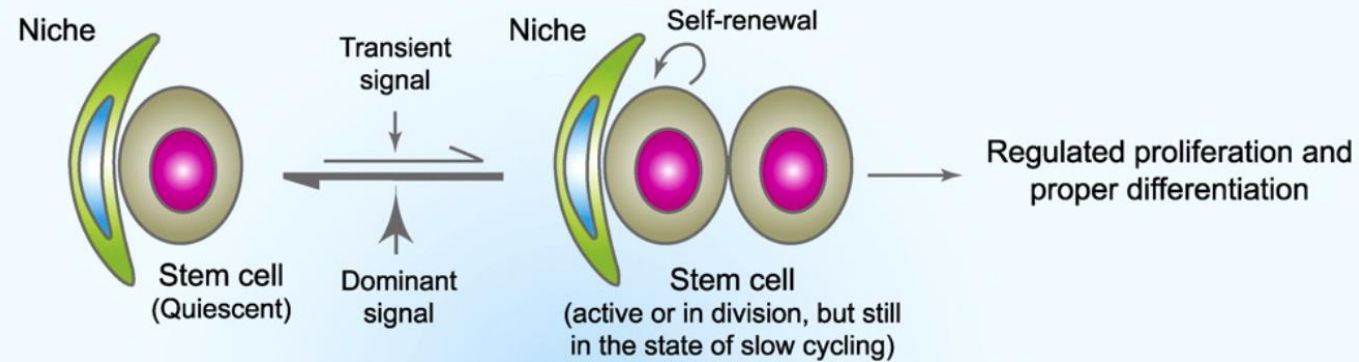
External control: role of microenvironment in haematopoiesis

- Produce growth factors
- Support and regulate balanced self-renewal and differentiation
- To increase mature cells in response to increased demand
- Physical support
- Homing of stem and progenitor cells

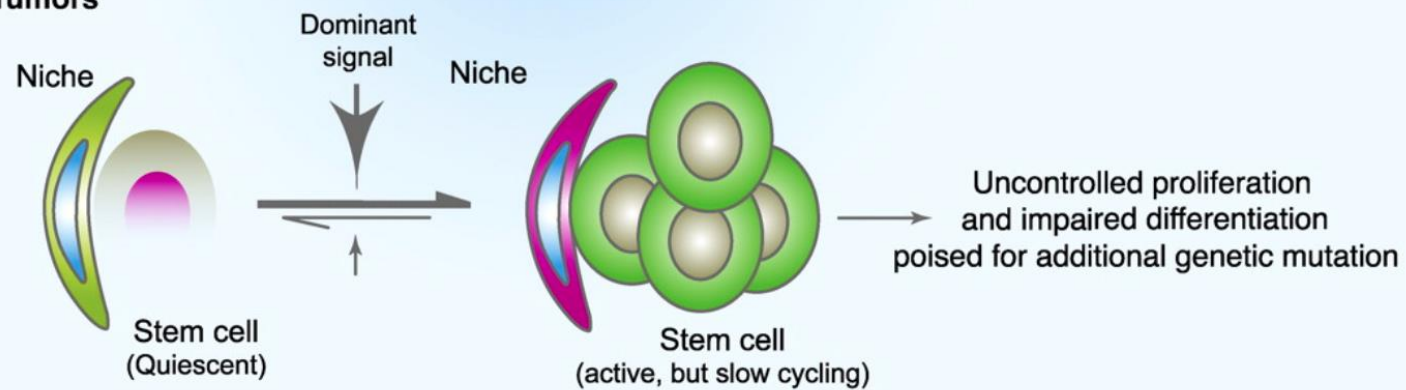


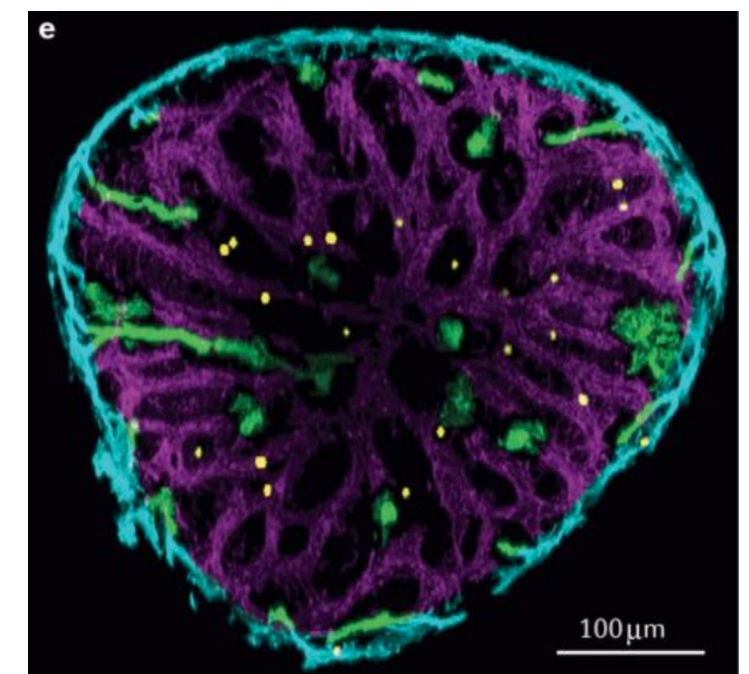
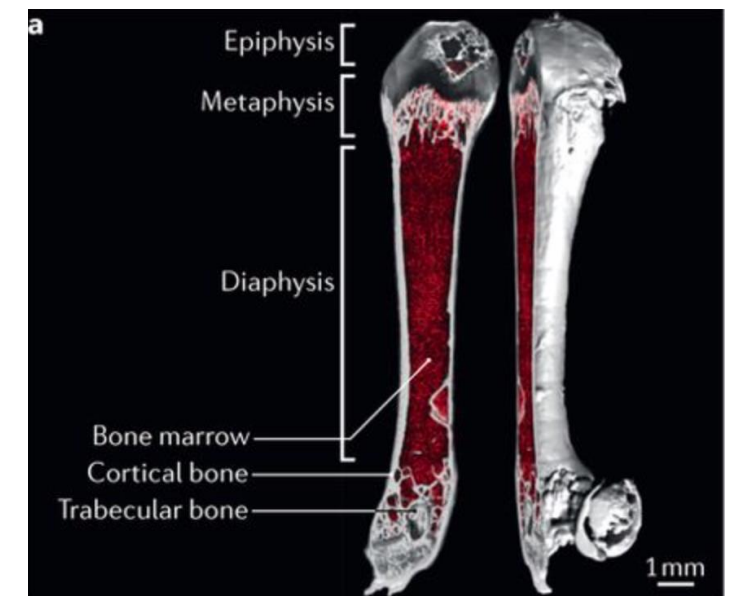
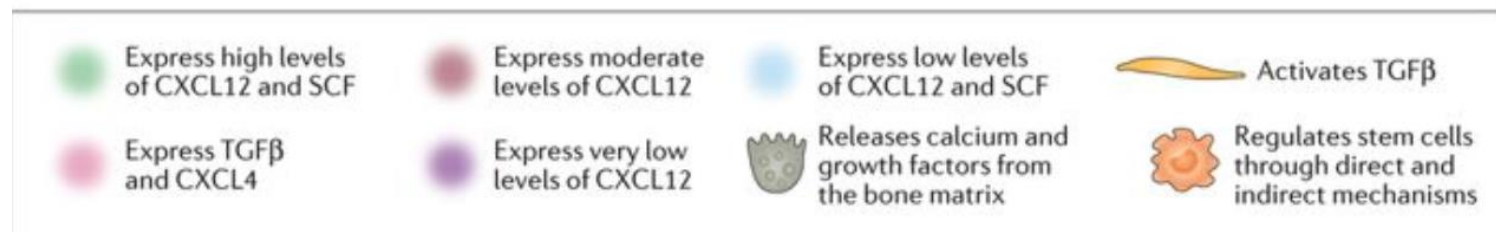
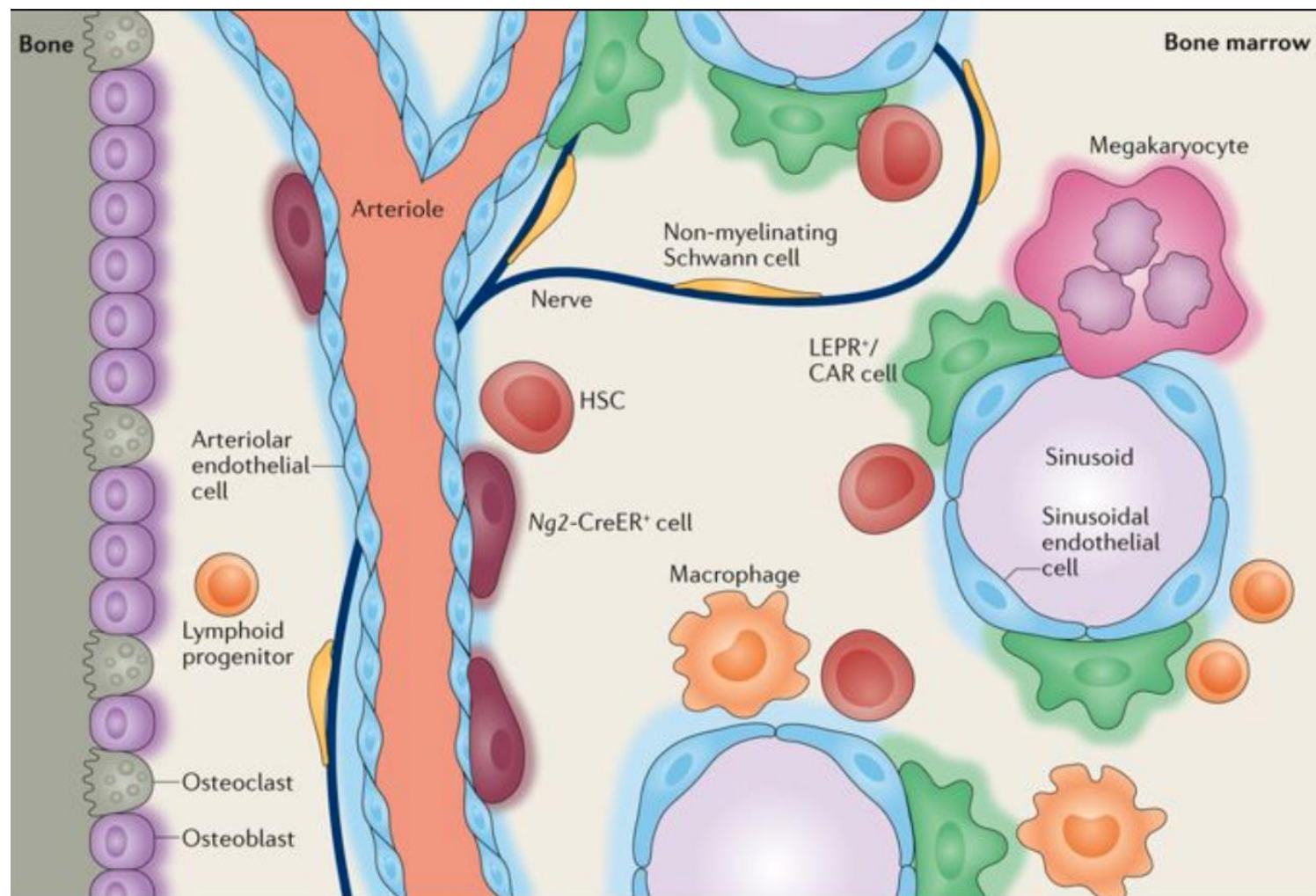
Niche-HSC interactions

Under Normal Physiological Conditions



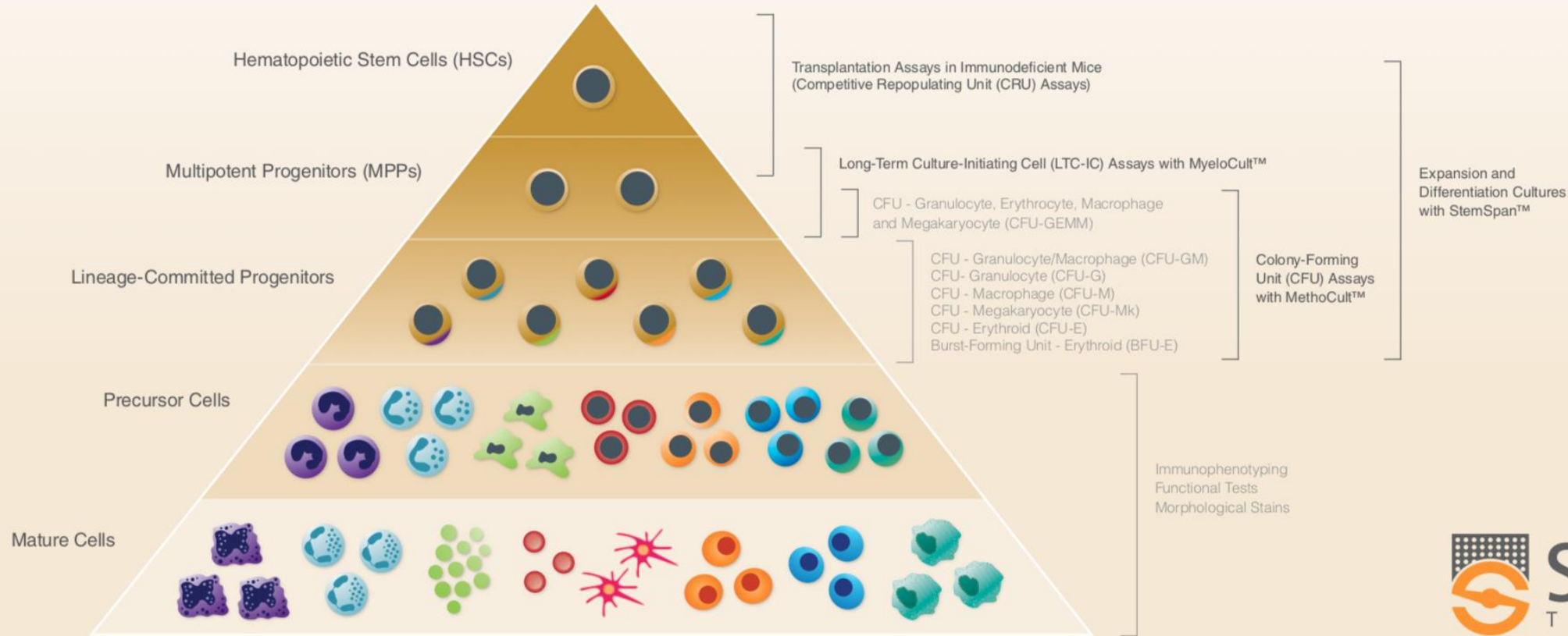
In Cancers or Tumors





Experimental methods to study HSCs

Hematopoietic Cell Compartments and Assays for Their Identification



References

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Experimental methods in HSC biology



A neon sign with the words "QUIZ TIME" in a stylized, outlined font. The sign is composed of two colors: bright pink and cyan. The word "QUIZ" is in pink, and "TIME" is in cyan. The sign is framed by a neon border that is also split between pink and cyan. The background is a dark blue brick wall.

QUIZ
TIME

Experimental methods in HSC biology

Which of the following is a 'gold standard' **stem cell** assay?

- (1) Single cell RNA-sequencing
- (2) Single cell colony forming assay
- (3) Injection of a single cell into irradiated mouse and determination of lineage read out
- (4) Single cell ATAC-sequencing

Experimental methods in HSC biology

- Which of the following cell surface markers are frequently used to identify multipotent stem/progenitor cells in humans?
 - (A) CD34
 - (B) Sca1
 - (C) cKIT
 - (D) CD38
 - (E) CD90

Experimental methods in HSC biology

Which of the following sentences is *not* true

- (A) Haematopoiesis is first observed in the AGM region around 5 weeks gestation
- (B) The bone marrow is the primary site of haematopoiesis from the 2nd trimester
- (C) Haematopoiesis can be observed in the spleen, liver and skin in certain types of blood malignancies
- (D) Platelets are produced in the lung

QUIZ
TIME

Experimental methods to study HSCs

- Stem cell assays to confirm ***multipotency***
 - Single cell transplantation of human cells into immunodeficient mice (serial transplantation)
 - Long term culture-initiating cell (LTC-IC) assays
 - Lineage tracing and barcoding approaches
- Methods to **determine lineage potential** of stem and progenitor cells
 - What lineages can this cell produce?
 - Colony forming assays (e.g. Methocult)
 - Liquid culture assays (supportive media with expansion/differentiation cytokines)
- **‘Phenotypic’ assays**
 - Immunophenotyping / proteomics
 - Gene expression assays

Need for a single cell approach

Bulk analysis



Information is 'averaged'

How do you determine if a population is homogeneous?

Is any population truly homogeneous??

Unable to understand functional read-outs and to properly compare two populations using 'bulk' assays

Need for a single cell approach

Bulk analysis



Information is 'averaged'

Single cell analysis



Unveil rare cell populations
Cellular hierarchies
Combinatorial patterns of gene expression

Need for a single cell approach

Bulk analysis



Single cell analysis



3D context/spatial positioning



Information is 'averaged'



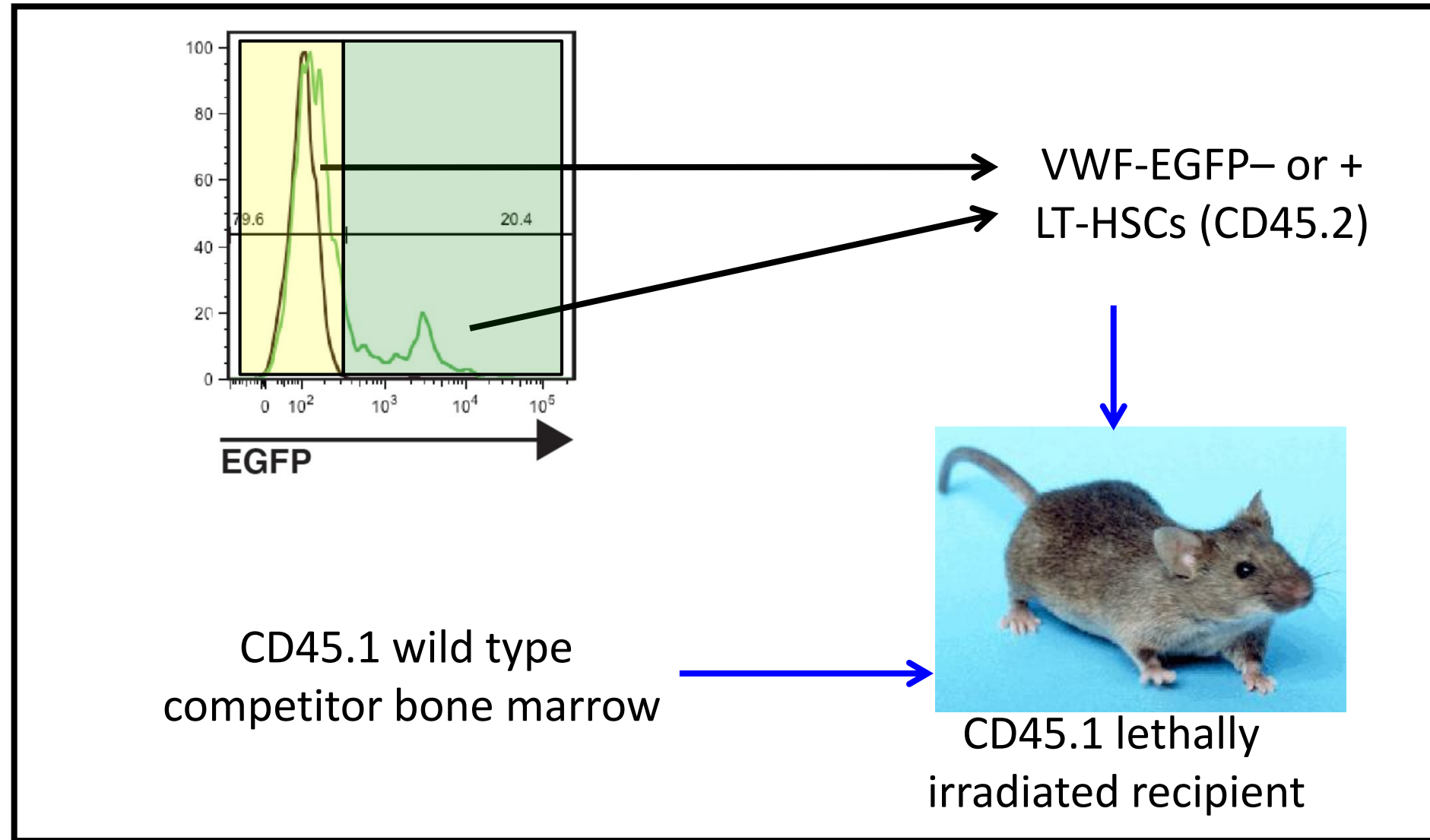
Unveil rare cell populations
Cellular hierarchies
Combinatorial patterns of gene expression

Organization in native context is crucial to understand cell-cell interactions

Single cell transplantation to identify lineage priming

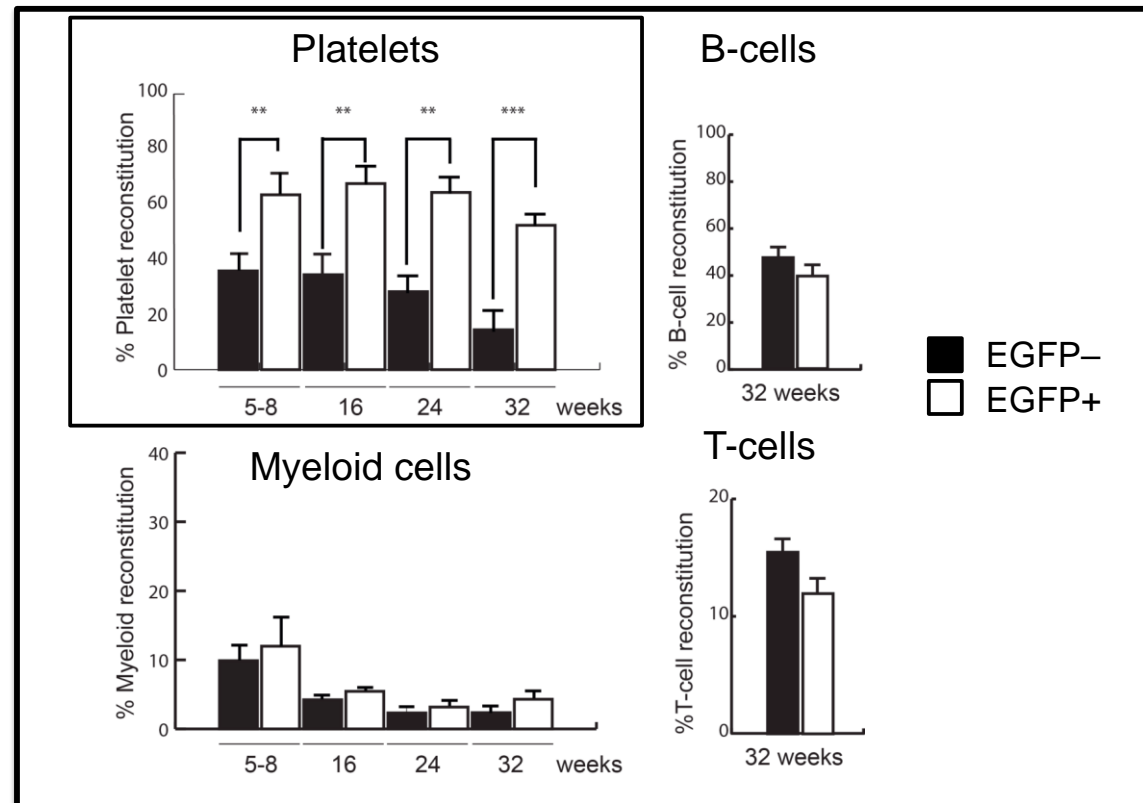
- 1st in vivo transplantation assay was developed in 1950s by Ford et al (1956) and McCulloch & Till 1960
- CFU-S: colony forming unit-spleen cells
 - Cells injected into an irradiated recipient formed macroscopic splenic colonies 1-3 weeks later
 - Not a stem cell assay
- **Long term repopulating assays**
 - Usually done as competitive repopulation assay, to measure a population with unknown repopulating ability against competitor whole BM containing a known number of HSC → calculate the repopulating units
 - Frequency of HSC can be measured using a limiting dilution assay (dilutions of the test cells against constant number of competitors, measure the number of mice that are reconstituted)

Single cell transplantation to identify lineage priming



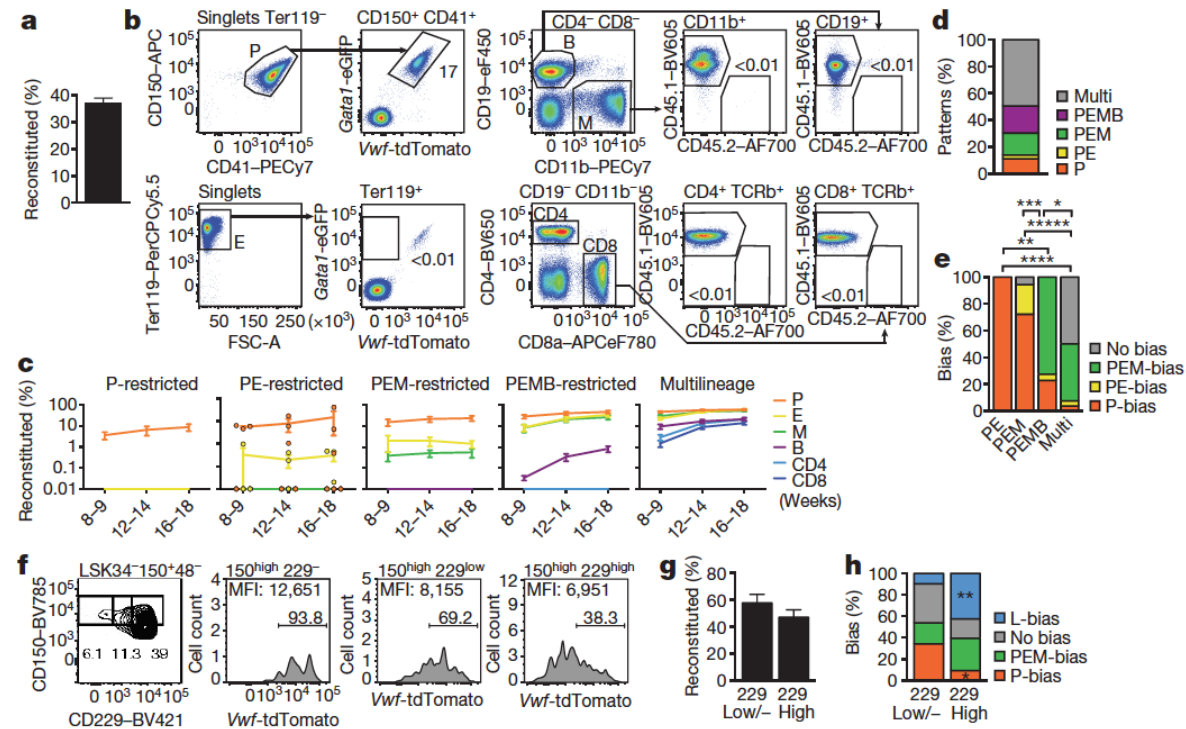
Platelet-biased stem cells reside at the apex of the haematopoietic stem-cell hierarchy

Alejandra Sanjuan-Pla^{1,2}, Iain C. Macaulay^{3,4}, Christina T. Jensen^{3,4}, Petter S. Woll^{3,4}, Tiago C. Luis^{3,4}, Adam Mead^{3,4}, Susan Moore^{1,2}, Cintia Carella², Sahoko Matsuoka^{3,4}, Tiphaine Bouriez Jones^{3,4}, Onima Chowdhury^{3,4}, Laura Stenson^{3,4}, Michael Lutteropp^{3,4}, Joanna C. A. Green^{3,4}, Raffaella Facchini^{3,4}, Hanane Boukarabila^{3,4}, Amit Grover³, Adriana Gambardella³, Supat Thongjuea³, Joana Carrelha^{3,4}, Paul Tarrant^{3,4}, Deborah Atkinson^{3,4}, Sally-Ann Clark^{3,4}, Claus Nerlov^{1,2,3*} & Sten Eirik W. Jacobsen^{3,4*}



Hierarchically related lineage-restricted fates of multipotent haematopoietic stem cells

Joana Carrelha^{1,2}, Yiran Meng^{1,2}, Laura M. Kettyle^{3,4}, Tiago C. Luis^{1,2}, Ruggiero Norfo^{1,2}, Verónica Alcolea^{1,2}, Hanane Boukarabila^{1,2†}, Francesca Grasso^{4,5}, Adriana Gambardella², Amit Grover², Kari Högstrand^{3,4}, Allegra M. Lord^{3,4}, Alejandra Sanjuan-Pla^{2‡}, Petter S. Woll^{4,5}, Claus Nerlov^{2*} & Sten Eirik W. Jacobsen^{1,2,3,4,5*}

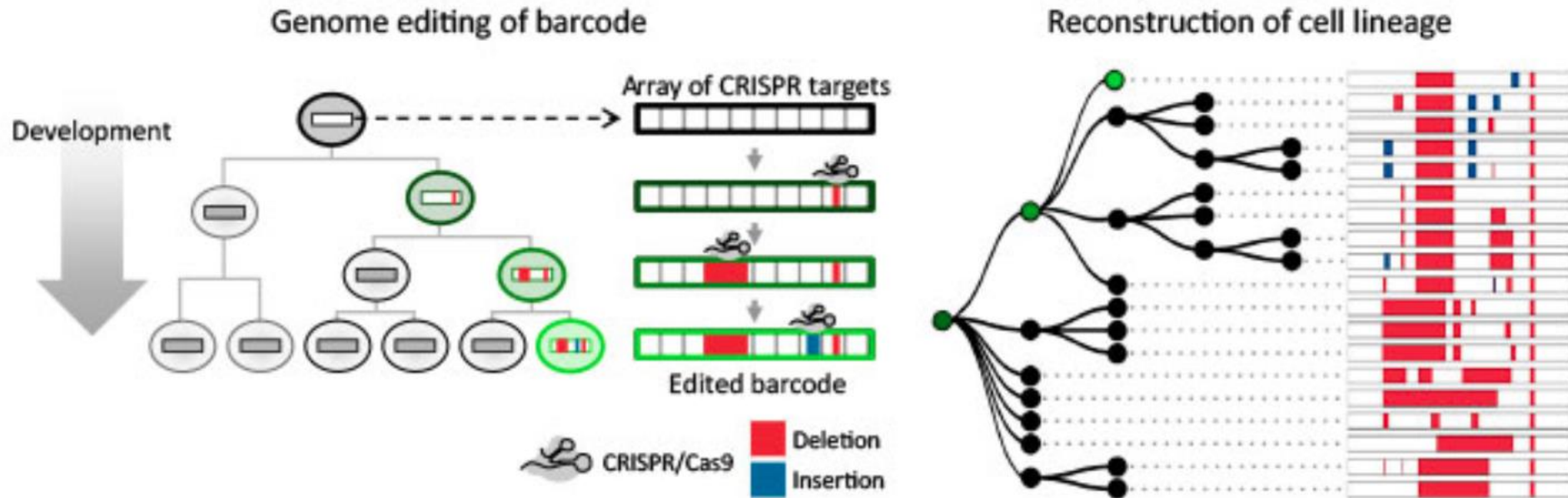


Considerations in these expts

- Cut off for engraftment is arbitrary – 1% for all lineages?
- Time point for analysis?
- Differences in genetic background of mice
- What method has been used to identify the donor vs recipient cells and how robust is this
- Can be difficult to determine whether a phenotype is due to differences in the HSC or in their potential to produce progeny
- Expensive and use a lot of mice!
- Assay also dependent on homing and engraftment potential of HSCs
- BM microenvironment is perturbed
- Not all lineages can be thoroughly assessed e.g. Mk-E

Lineage tracing / barcoding assays

- CRISPR-Cas9: ‘dropping genetic breadcrumbs’ into developing cells



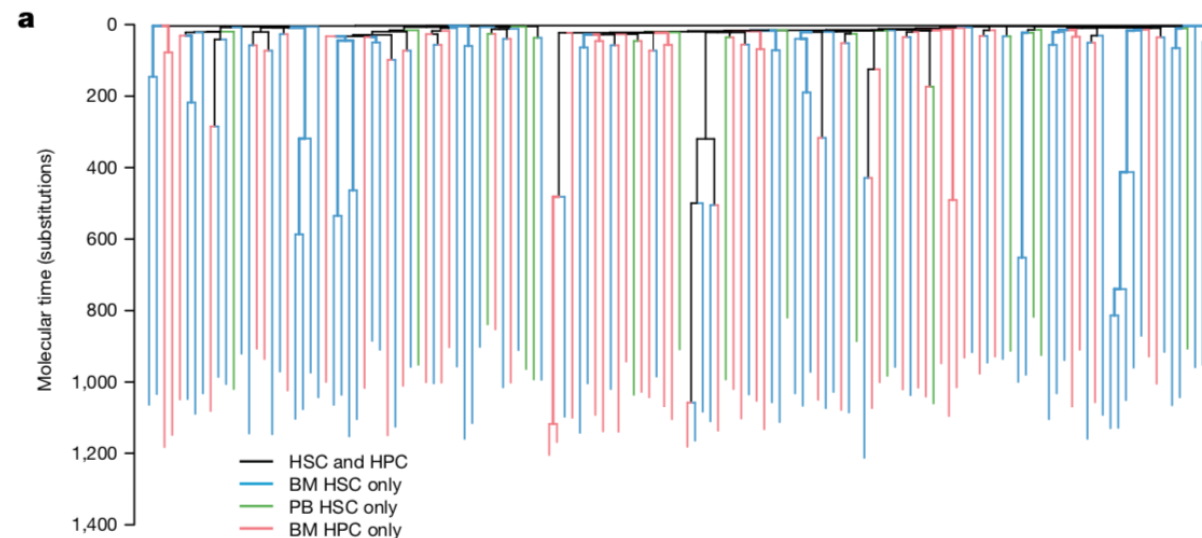
GESTALT – Science 2016. CRISPR-Cas9 used to introduce a 300 bp ‘barcode’ into dividing cells and cell progeny then tracked as they accumulate mutations. Progeny cells then collected and DNA amplified to ‘read’ the barcode and reconstruct the lineage tree

ARTICLE

<https://doi.org/10.1038/s41586-018-0497-0>

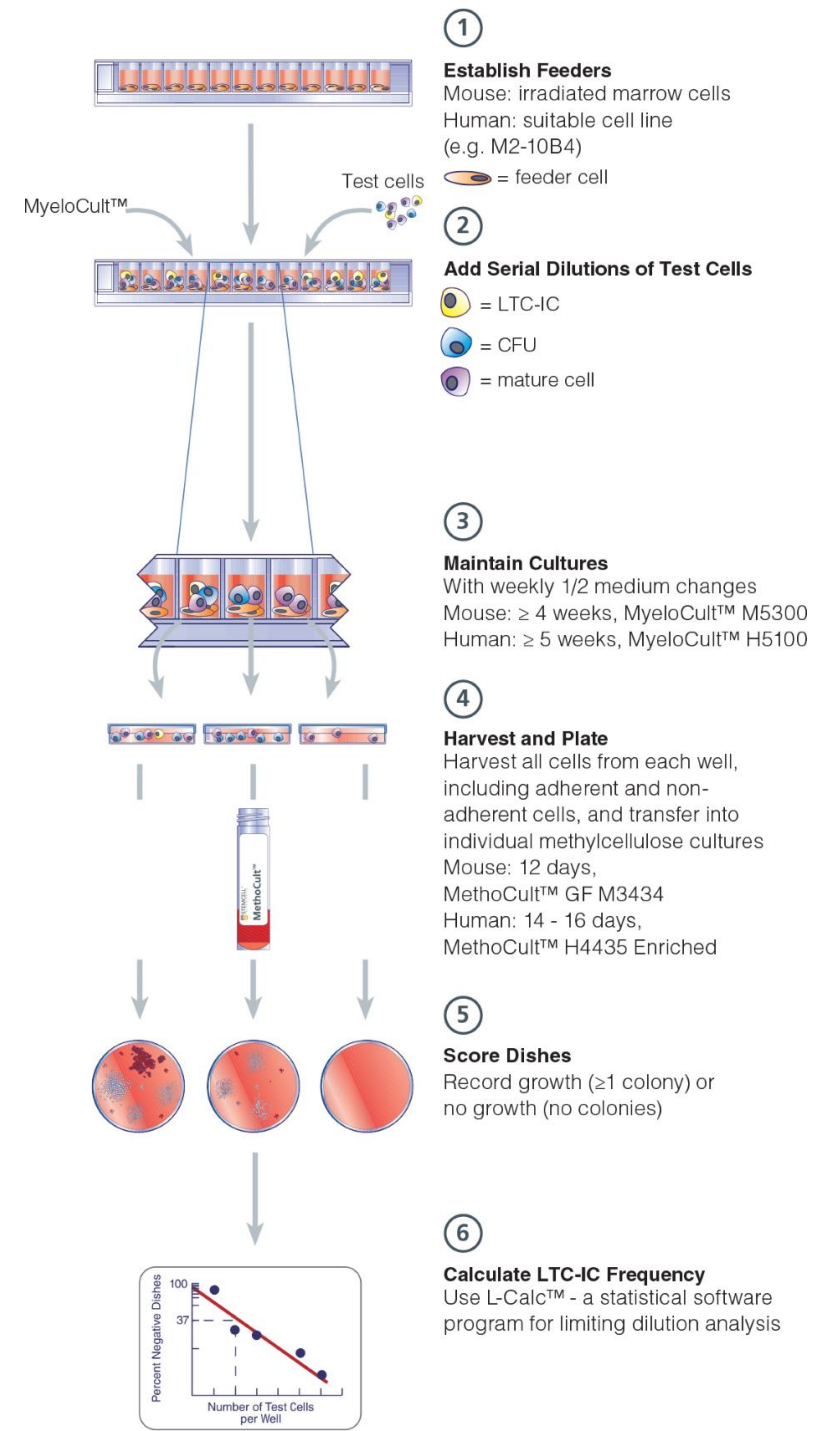
Population dynamics of normal human blood inferred from somatic mutations

Henry Lee-Six¹, Nina Friesgaard Øbro², Mairi S. Shepherd², Sebastian Grossmann¹, Kevin Dawson¹, Miriam Belmonte², Robert J. Osborne¹, Brian J. P. Huntly², Inigo Martincorena¹, Elizabeth Anderson¹, Laura O'Neill¹, Michael R. Stratton¹, Elisa Laurenti², Anthony R. Green^{2,3*}, David G. Kent^{2,3*} & Peter J. Campbell^{1,3*}



In vitro stem cell assay: Long term Colony-initiating cell assay (LTC-IC)

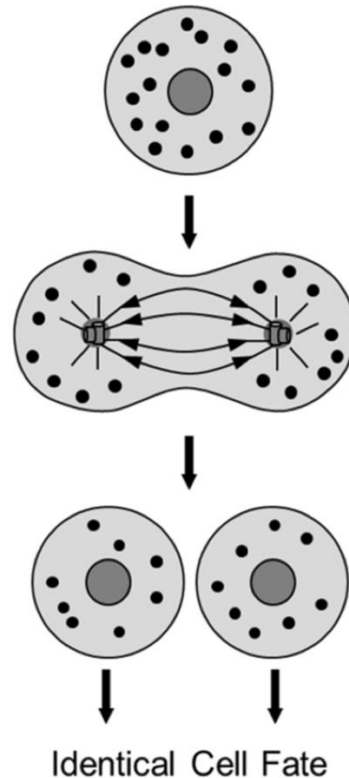
Plate cells on stromal support cells for 4-6 weeks
Then transfer to methylcellulose to see if can initiate colonies



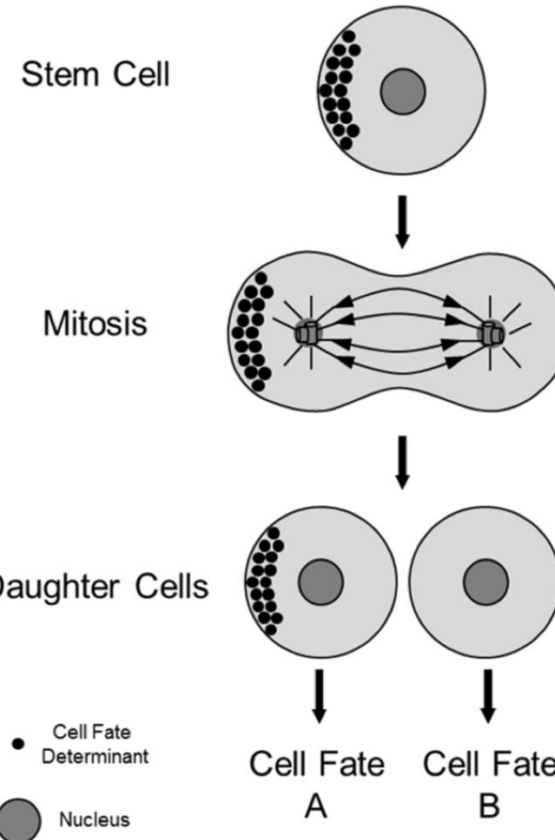
Paired daughter cell assays

Micromanipulation of progeny following 1st cell division to test symmetric vs asymmetric division

Symmetric cell division



Asymmetric cell division

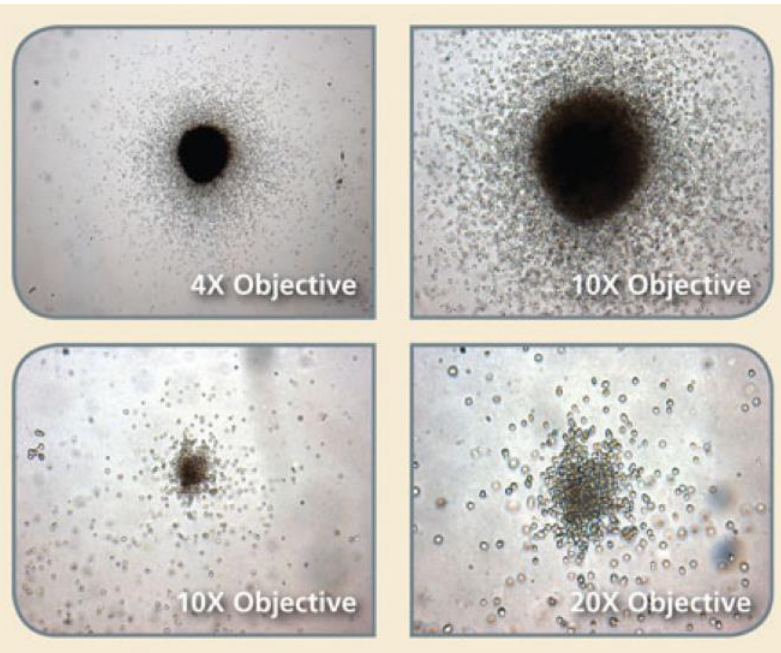


In vitro differentiation assays

Semi-solid 'clonogenic' assays: mix cells with methylcellulose (+ cytokines)

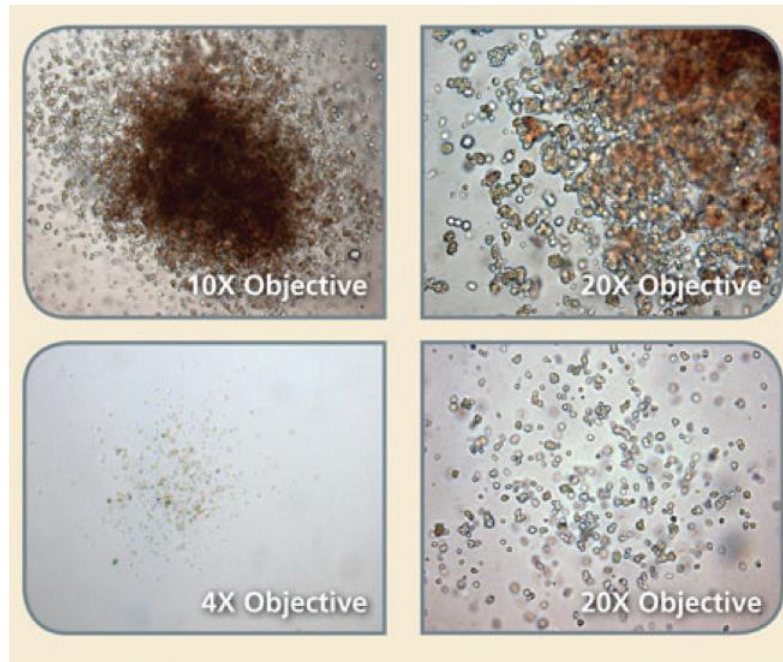
CFU-GM

Ery



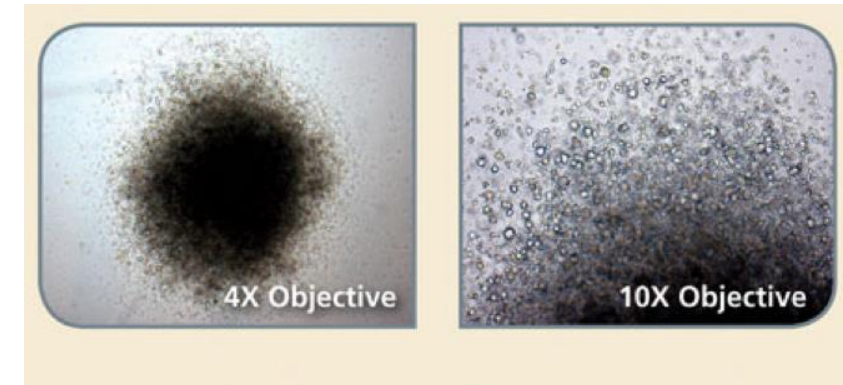
BFU-E

Ery



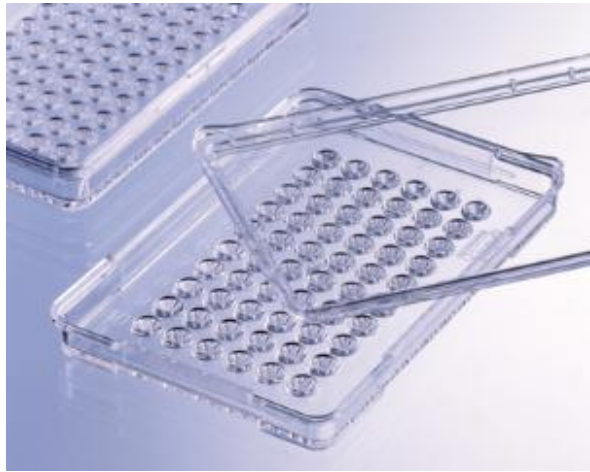
CFU-GEMM

Gran, ery, macrophage, mega



In vitro differentiation assays

- Liquid culture (single cell 96-well plate or Terasaki plates)



Stemspan (serum free media)

- + Stem cell Factor
- + erythropoietin
- + thrombopoietin
- + G-CSF, GM-CSF
- + Interleukins IL3, 6
- + FLT3-Ligand

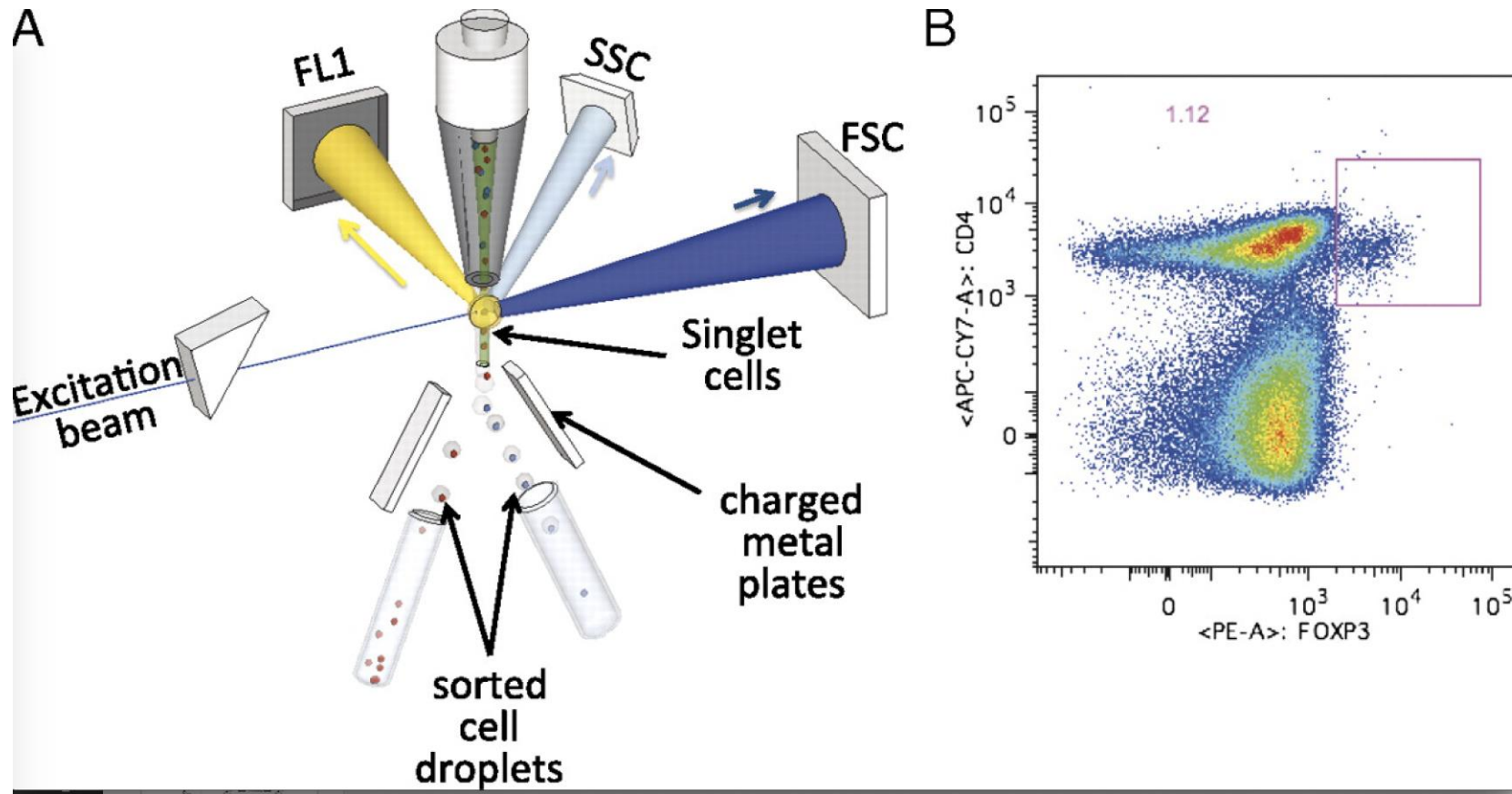
May need stromal support cells

- OP9 for B –ells
- OP9DL1 for T cells

Semi-solid / liquid culture pitfalls

- Measure progenitor frequency rather than HSCs
- Assessing multipotency of single cells is really challenging
- Readout is very dependent on skills and experience of observer!
- Biased by culture conditions / cytokines added

Phenotypic assays: immunophenotyping



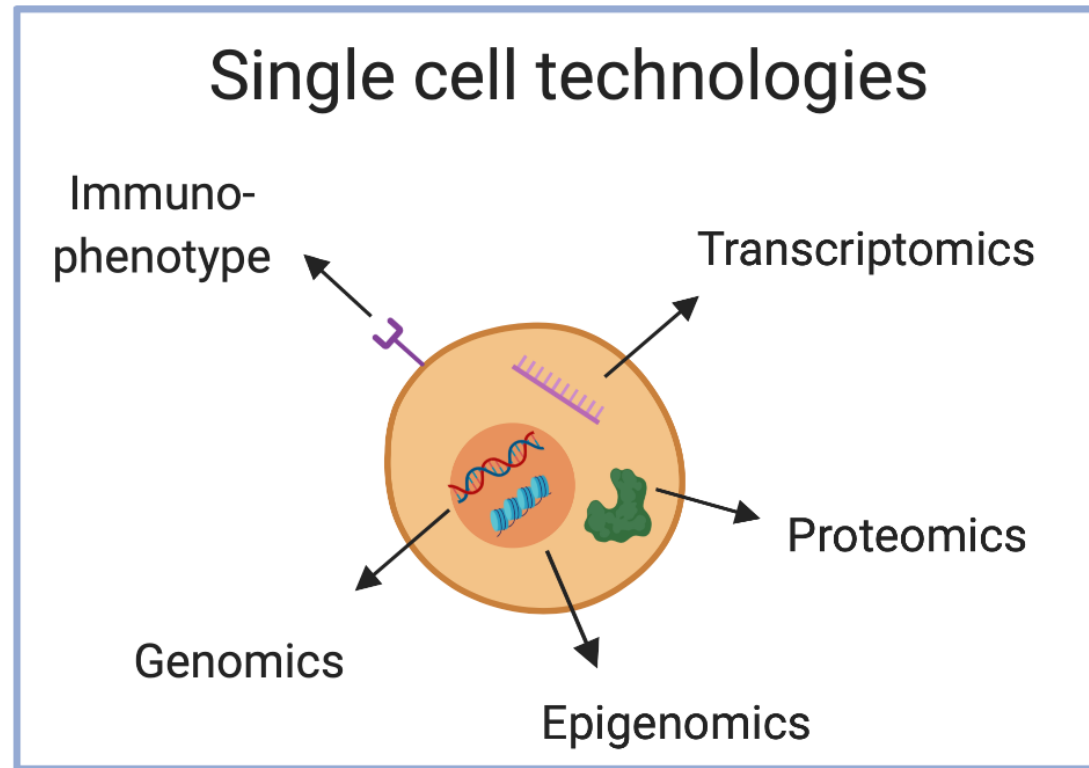
Cells are excited by laser beams, to delineate forward scatter (FSC) and side scatter (SSC) and labelled with fluorescently-labelled antibodies detected by fluorescent light (FL1)

Jaye et al, J of Immunol 2012

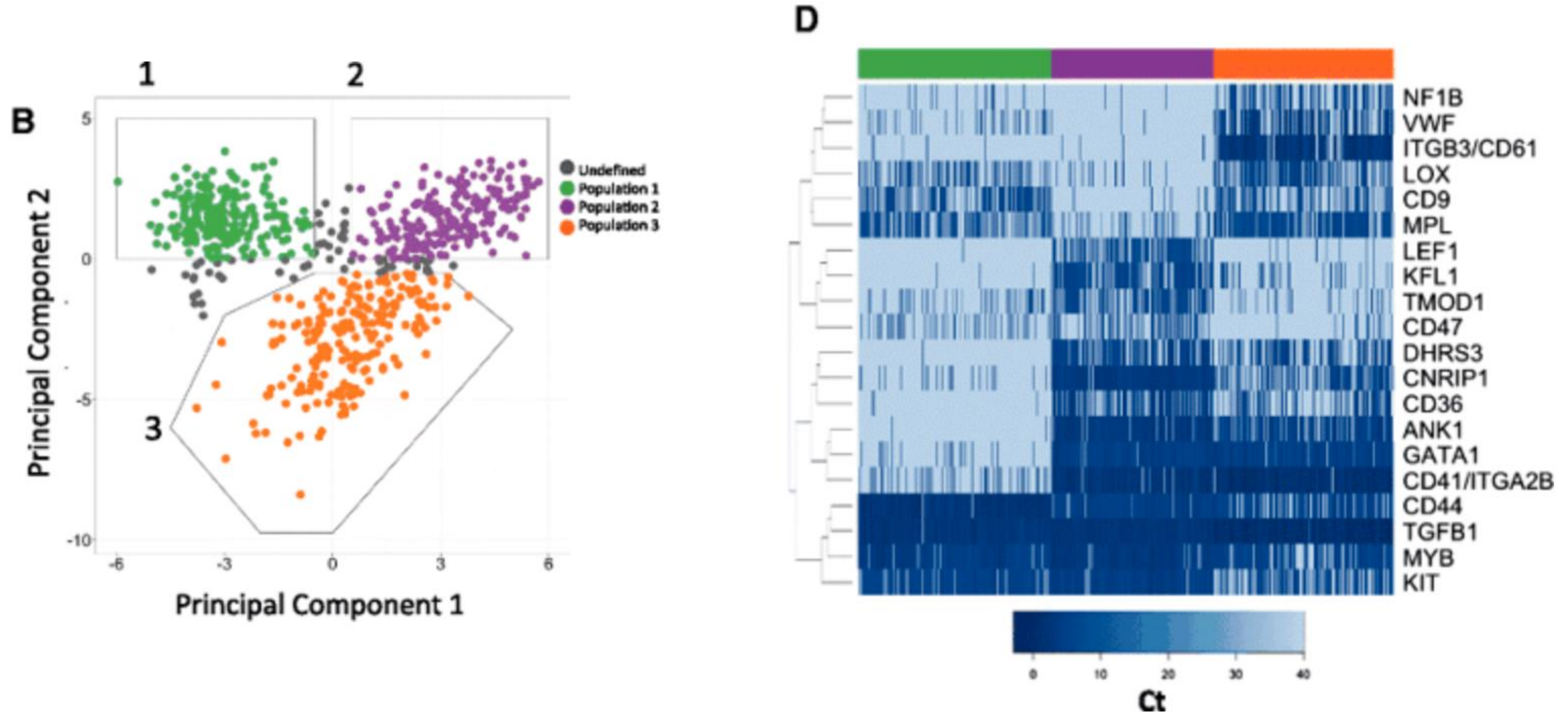
Phenotypic assays: immunophenotyping

- Various different immunophenotyping strategies for mouse and human
 - Purity of HSCs identified using the most state-of-the-art FACS methods is thought to be around 50% (assessed by the transplantation assay)
 - Variability in HSC antigens between different mouse strains
 - Different stages of ontogeny
 - Cell cycle
 - Non-steady state e.g. post transplant/5-FU treatment
- 'true' HSC need to be defined by functional analyses

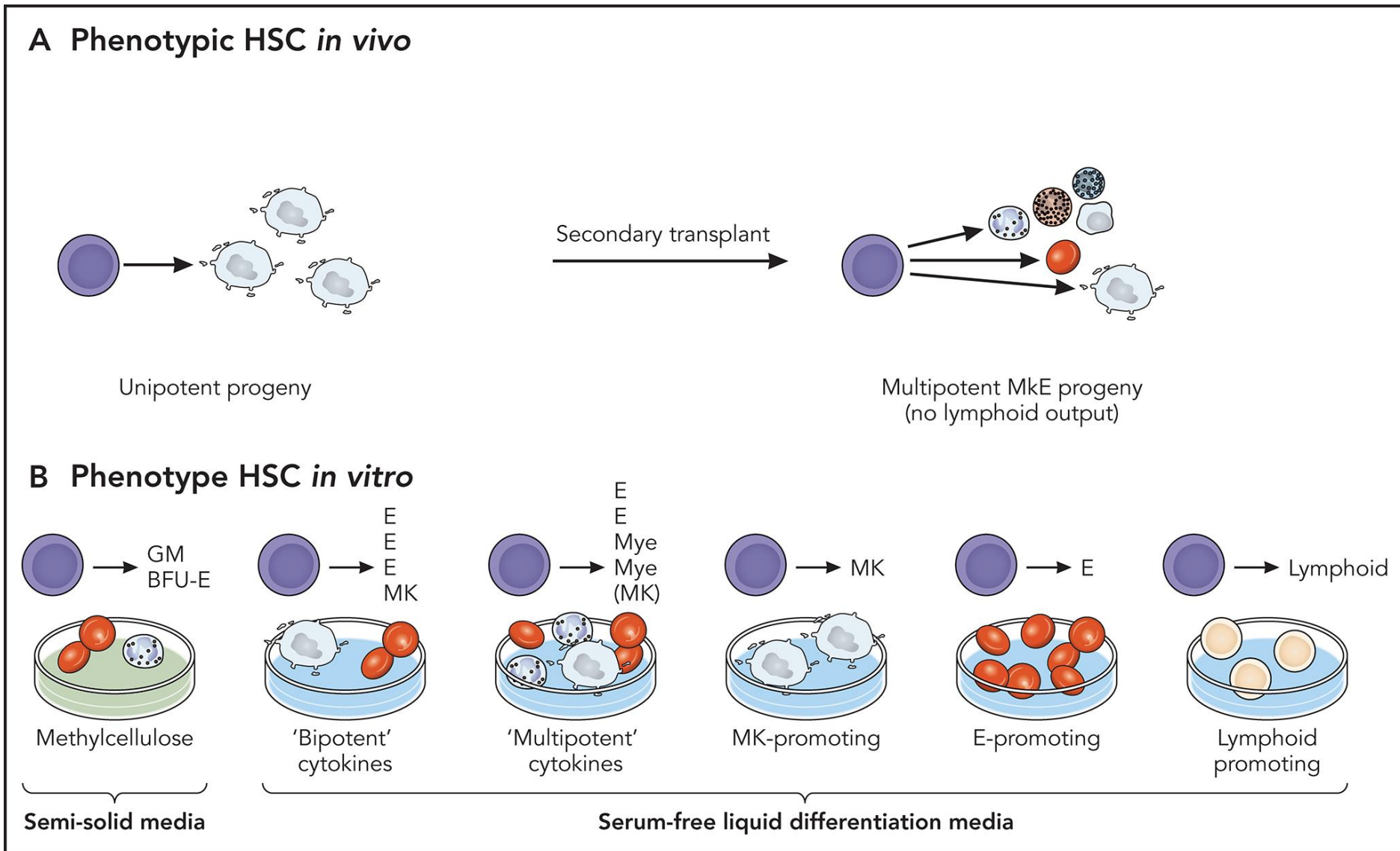
Molecular profiling to identify stem/progenitor priming



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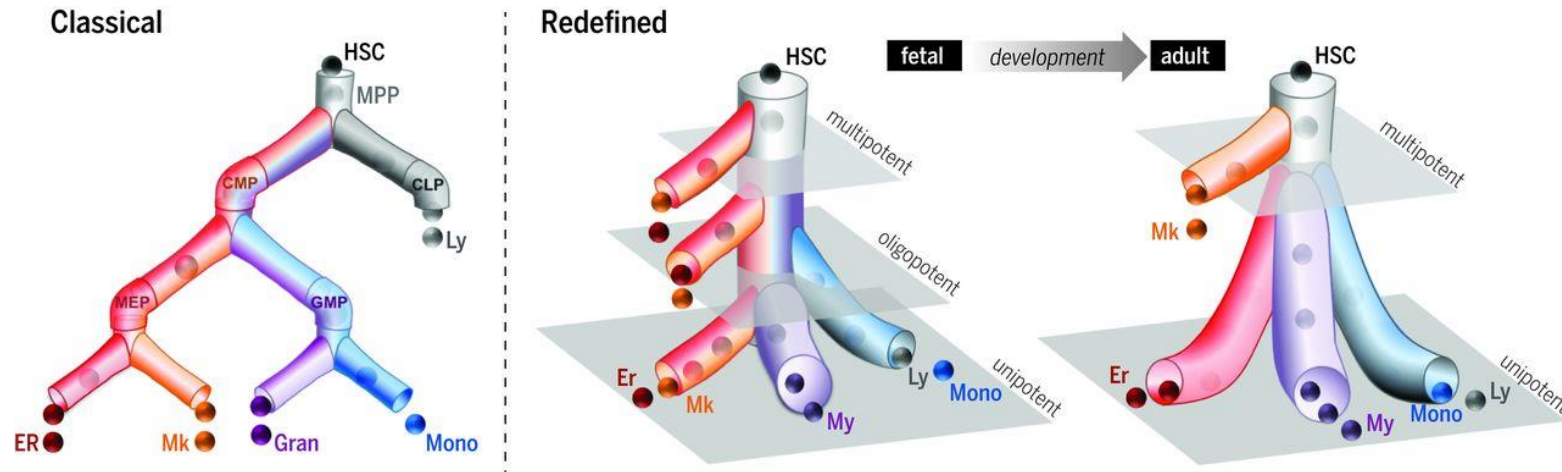


Limitations in stem cells assay even at single cell level



Questions/controversies

- Does haematopoiesis occur via step-wise transitions or over a gradual continuum of differentiation?
- What % of HSCs are truly 'multipotent'? Recent data suggests that majority of HSCs in adult life are 'biased' towards certain differentiation fates



Questions/controversies

- Do HSCs contribute to steady-state haematopoiesis or are they a 'reserve' (with most contributions coming from progenitors)?
- How much similarity is there between humans and animal models?

Why does this matter?

Abnormalities of blood cell production

- **Too few cells produced**

- Bone marrow failure e.g. aplastic anaemia
- Anaemia, neutropenia, thrombocytopenia

- **Too many cells produced**

- Essential thrombocythaemia
- Polycythaemia vera
- Chronic myeloid leukaemia

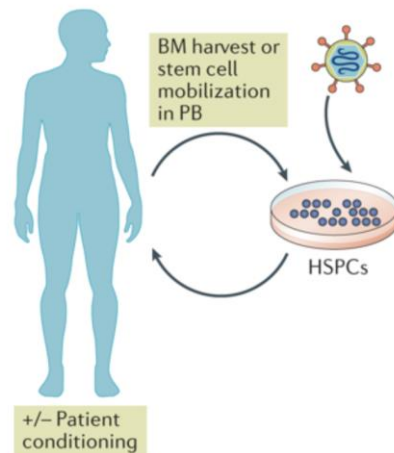
- **Abnormal cell production**

- Malignant: leukaemia, myelodysplasia
- Non-malignant: sickle-cell anaemia

What are the potential clinical uses of HSCs?

What are the potential clinical uses of HSCs?

- Transplantation for haematopoietic disorders e.g. leukaemia
- In vitro production of red cells, platelets
- Repair of damaged tissue e.g. heart, liver
- Gene therapy e.g. SCID, thalassaemia



See review by Cavazzana et al, Nature Reviews 2019

Summary

- Haematopoiesis migrates from YS --> AGM --> FL → FBM in fetal life
 - Postnatal life, continues in bone marrow
 - Pathological extramedullary haematopoiesis can occur in liver, spleen, almost anywhere!
 - Regulated by key genes and transcription factors
 - Regulated by growth factors and the microenvironment
 - Review of *in vivo* and *in vitro* experimental techniques and challenges
- need to carefully evaluate what has been done and what the assay is ***actually*** testing (cell fate vs potential) to interpret the literature

Questions?