

Modelling hematopoietic stem cells and their interaction with the bone marrow micro-environment.

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Stem Cell Modelling Day
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- ▶ Brief introduction to mathematical modelling.

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- ▶ Brief introduction to stem cells.

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- ▶ Brief introduction to stem cells.
- ▶ Development of a mathematical model.

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- ▶ Brief introduction to mathematical modelling.
- ▶ Brief introduction to stem cells.
- ▶ Development of a mathematical model.
- ▶ What does the model tell us?

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- ▶ A way to test hypotheses

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- ▶ A way to test hypotheses
- ▶ “What if the world worked like this?”

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- ▶ A way to test hypotheses
- ▶ “What if the world worked like this?”
- ▶ Does it agree with intuition?

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- ▶ A way to test hypotheses
- ▶ “What if the world worked like this?”
- ▶ Does it agree with intuition?
- ▶ With biological theories?

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- ▶ A way to test hypotheses
- ▶ “What if the world worked like this?”
- ▶ Does it agree with intuition?
- ▶ With biological theories?
- ▶ With data?

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Stem cells (As seen by a mathematician)

- ▶ “Stem cells are cells that can differentiate into other types of cells, and can also divide in self-renewal to produce more of the same type of stem cells.” - Wikipedia intro.

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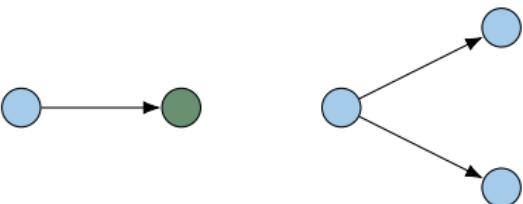
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Stem cells (As seen by a mathematician)

- ▶ “Stem cells are cells that can differentiate into other types of cells, and can also divide in self-renewal to produce more of the same type of stem cells.” - Wikipedia intro.



where are stem cells and is a differentiated/progenitor cell.

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Stem cells (As seen by a mathematician)

- ▶ “Stem cells are cells that can differentiate into other types of cells, and can also divide in self-renewal to produce more of the same type of stem cells.” - Wikipedia intro.
- ▶ Very hard to measure *in vivo*.

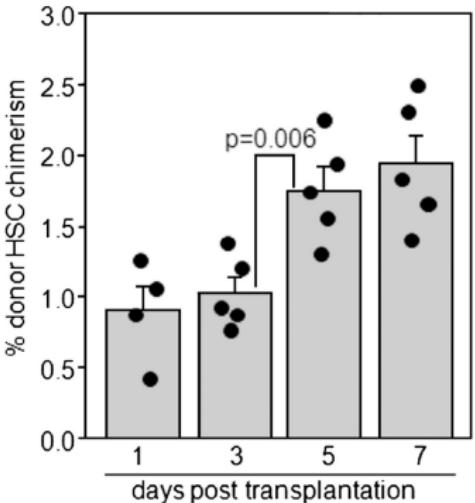


Figure: Mouse experiment data (Bhattacharya et al., 2009)

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- ▶ *Cancitis* group at RUC: Modelling of development and treatment of blood cancers (leukemias), in particular Myeloproliferative Neoplasms (MPNs).

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- ▶ *Cancitis* group at RUC: Modelling of development and treatment of blood cancers (leukemias), in particular Myeloproliferative Neoplasms (MPNs).
- ▶ Hematopoietic stem cells (HSCs) give rise to a vast production of blood cells.

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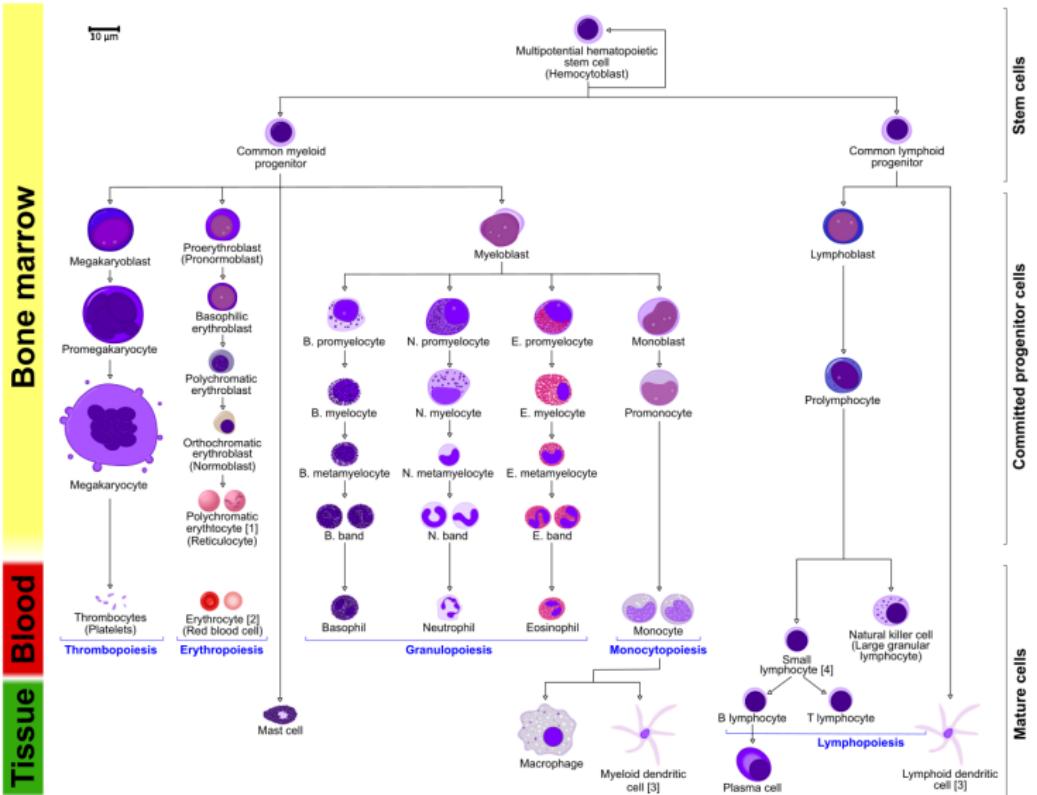
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- ▶ Hematopoietic stem cells (HSCs) give rise to a vast production of blood cells.
- ▶ Mutations of HSCs are believed to be central in the development of most leukemias.

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- ▶ Stem cell “niches” in the bone marrow micro-environment.

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- ▶ Mutations of HSCs are believed to be central in the development of most leukemias.
- ▶ Stem cell “niches” in the bone marrow micro-environment.

(Ashcroft et al., 2017), (Wang, Stiehl et al. 2017),
(Becker et al., 2019), (Wilson and Trumpp, 2006).

Modelling of the HSCs and their niches

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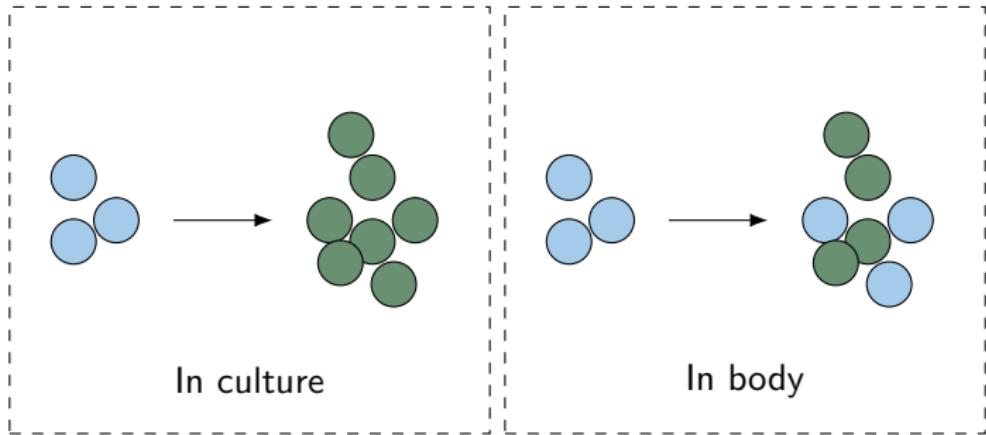
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Central hypothesis: Limited division, exhaustion after division.

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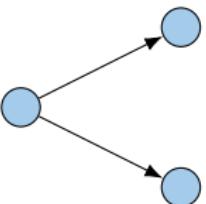
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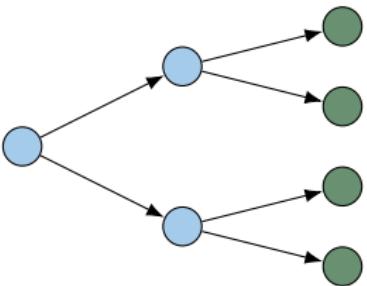
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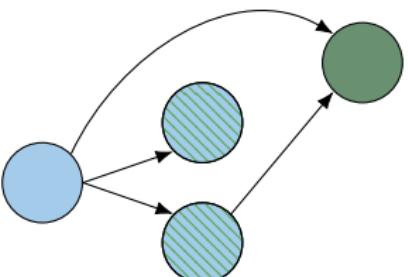
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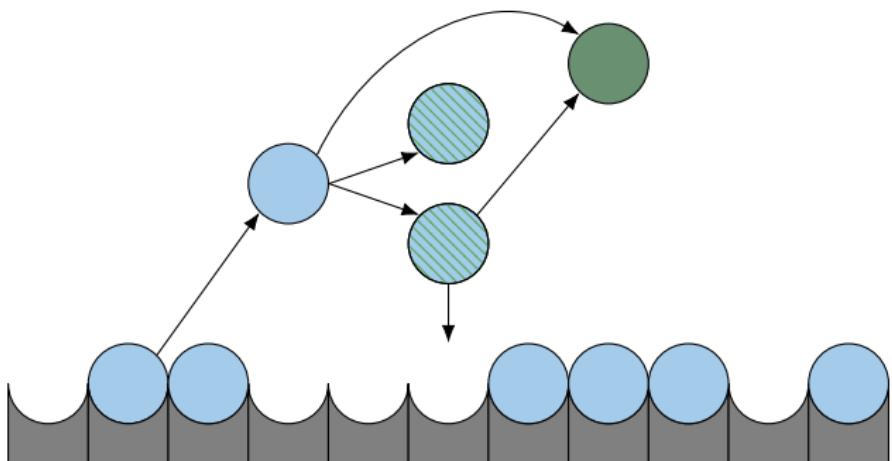
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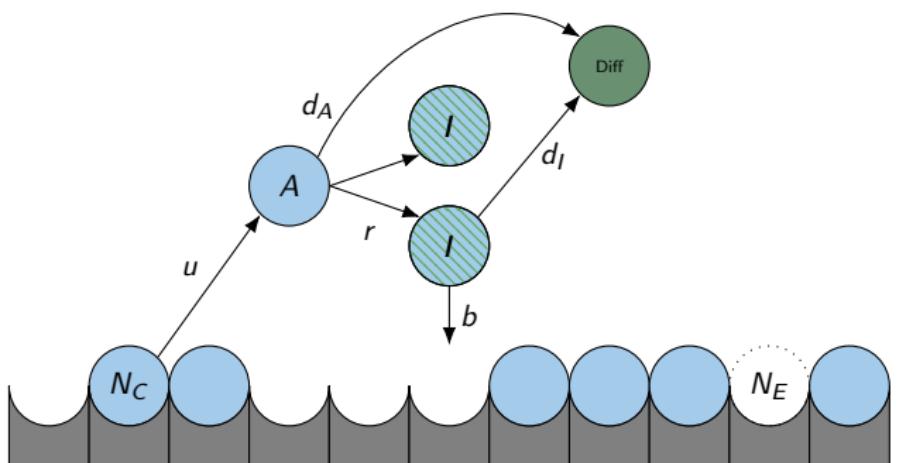
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N_C : Niche-bound, A : Active, I : Inhibited, N_E : Empty niches

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Central hypothesis: Limited division, exhaustion after division.

$$\frac{dN_E}{dt} = -bN_E I + uN_C$$

$$\frac{dN_C}{dt} = bN_E I - uN_C$$

$$\frac{dI}{dt} = -bN_E I + 2rA - d_I I$$

$$\frac{dA}{dt} = uN_C - rA - d_A A$$

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- ▶ The total number of niches (empty or cell-bound) are constant.

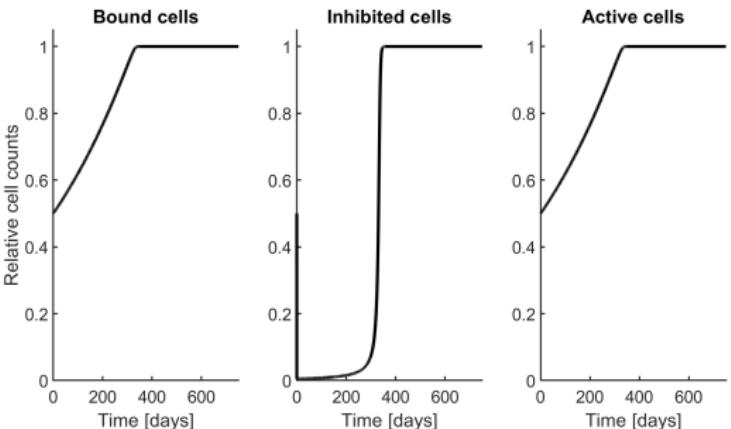
$$\frac{dN_E}{dt} + \frac{dN_C}{dt} = 0 \quad \Rightarrow \quad N_E + N_C = K$$

Initial analysis of the mathematical model

- The total number of niches (empty or cell-bound) are constant.

$$\frac{dN_E}{dt} + \frac{dN_C}{dt} = 0 \quad \Rightarrow \quad N_E + N_C = K$$

- It must be the case that active cells self-renew more than differentiate ($r > d_A$).

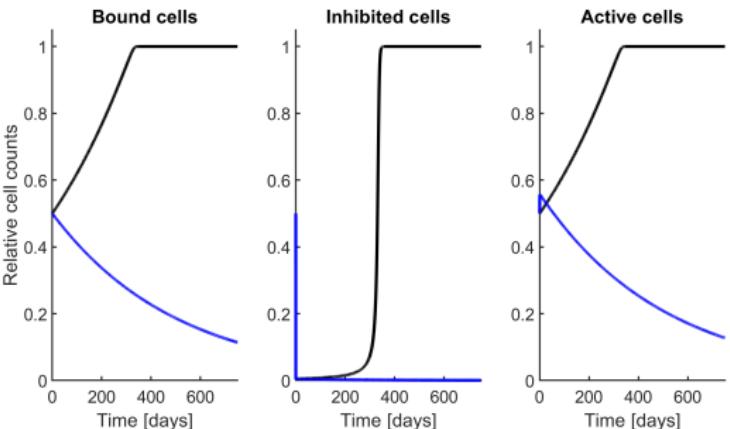


Initial analysis of the mathematical model

- The total number of niches (empty or cell-bound) are constant.

$$\frac{dN_E}{dt} + \frac{dN_C}{dt} = 0 \quad \Rightarrow \quad N_E + N_C = K$$

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- ▶ The total number of niches (empty or cell-bound) are constant.

$$\frac{dN_E}{dt} + \frac{dN_C}{dt} = 0 \quad \Rightarrow \quad N_E + N_C = K$$

- ▶ It must be the case that active cells self-renew more than differentiate ($r > d_A$).
- ▶ Number of empty niches in a “healthy” state is independent of unbinding from the niche (u).

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- ▶ The total number of niches (empty or cell-bound) are constant.

$$\frac{dN_E}{dt} + \frac{dN_C}{dt} = 0 \quad \Rightarrow \quad N_E + N_C = K$$

- ▶ It must be the case that active cells self-renew more than differentiate ($r > d_A$).
- ▶ Number of empty niches in a “healthy” state is independent of unbinding from the niche (u).
- ▶ If all niches disappear, cells die out.

Competition for niche-space

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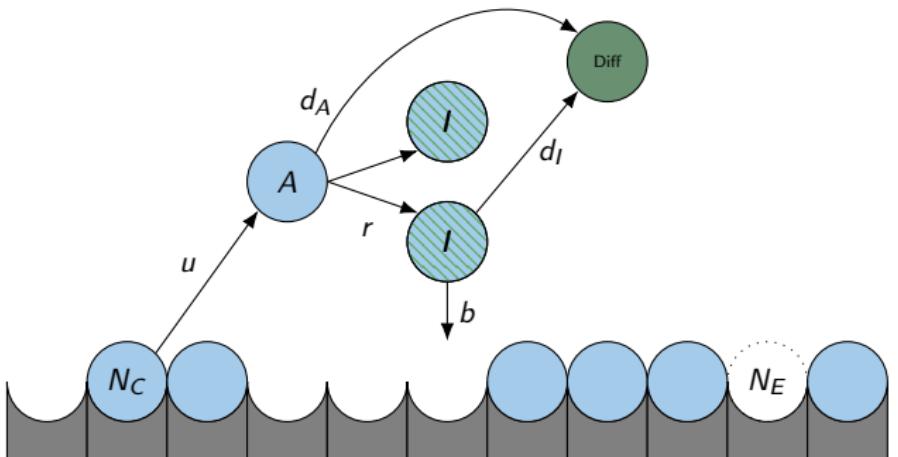
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N_C : Niche-bound, A : Active, I : Inhibited, N_E : Empty niches

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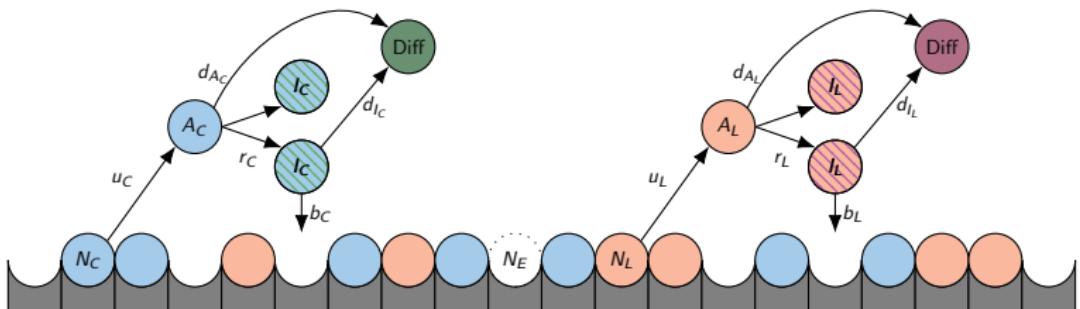
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N_C : Niche-bound C-type, A_C : Active C-type, I_C : Inhibited C-type

N_L : Niche-bound L-type, A_L : Active L-type, I_L : Inhibited L-type

N_E : Empty niches.

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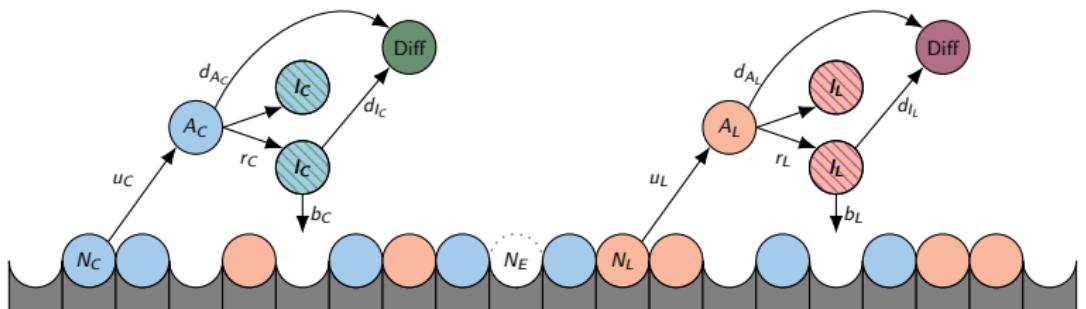
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Four steady states:

- ▶ No cell (Death)
- ▶ Only healthy cells (Hematopoiesis)
- ▶ Only leukemic cells (Full blown disease)
- ▶ Co-existence

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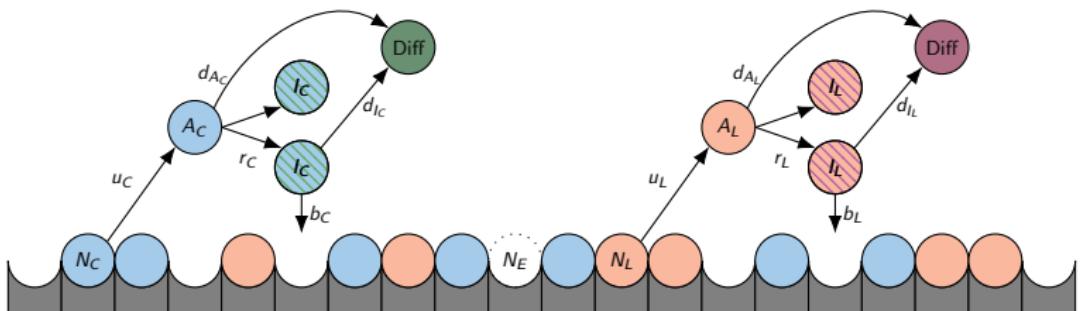
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Four steady states:

- ▶ No cell (Death)
- ▶ Only healthy cells (Hematopoiesis)
- ▶ Only leukemic cells (Full blown disease)
- ▶ Co-existence (Suppressed disease?)

Competition for niche-space - Fitness

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$$F_C = \frac{b_C}{d_{I_C}} \frac{(r_C - d_{A_C})}{(r_C + d_{A_C})} \quad \text{and} \quad F_L = \frac{b_L}{d_{I_L}} \frac{(r_L - d_{A_L})}{(r_L + d_{A_L})}$$

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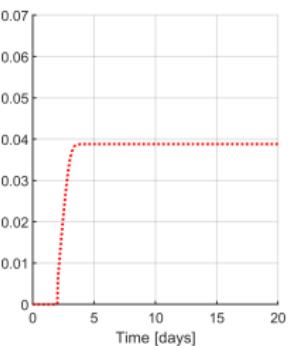
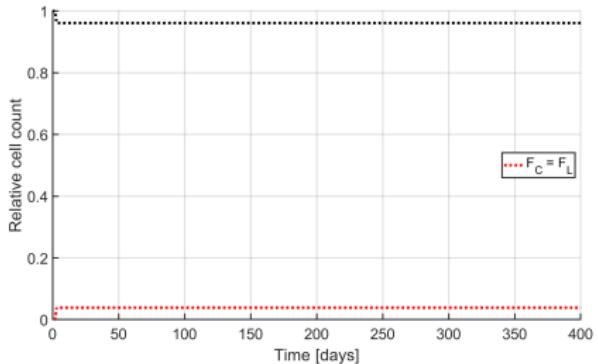
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$$F_C = \frac{b_C}{d_{I_C}} \frac{(r_C - d_{A_C})}{(r_C + d_{A_C})} \quad \text{and} \quad F_L = \frac{b_L}{d_{I_L}} \frac{(r_L - d_{A_L})}{(r_L + d_{A_L})}$$

- If $F_C = F_L$ then coexistence is possible.



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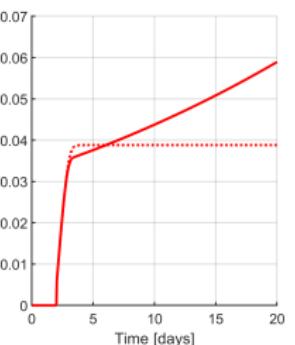
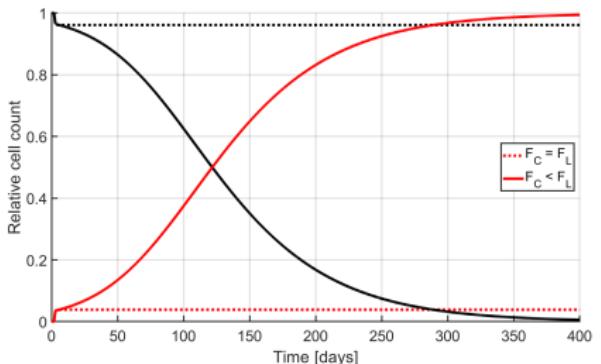
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- ▶ If $F_C = F_L$ then coexistence is possible.
- ▶ If $F_C < F_L$ then L -type outcompetes C -type



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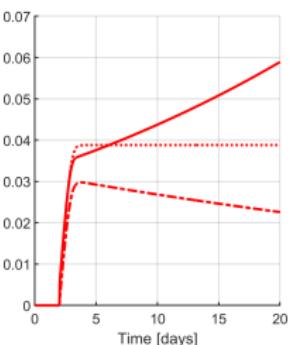
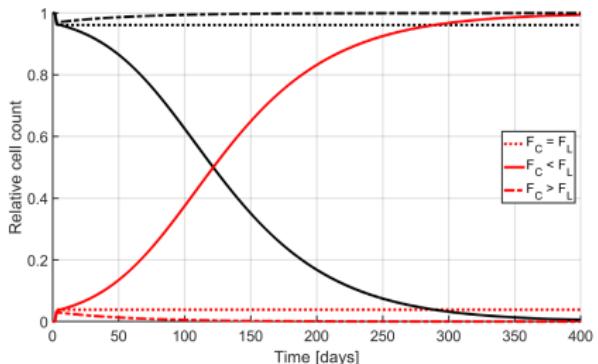
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$$F_C = \frac{b_C}{d_{I_C}} \frac{(r_C - d_{A_C})}{(r_C + d_{A_C})} \quad \text{and} \quad F_L = \frac{b_L}{d_{I_L}} \frac{(r_L - d_{A_L})}{(r_L + d_{A_L})}$$

- ▶ If $F_C = F_L$ then coexistence is possible.
- ▶ If $F_C < F_L$ then *L*-type outcompetes *C*-type
- ▶ If $F_C > F_L$ then *C*-type outcompetes *L*-type



Now what?

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- Model describing central mechanisms of HSCs.

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- ▶ Model describing central mechanisms of HSCs.
- ▶ Can be fit to experimental (mouse) data.

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- ▶ Model describing central mechanisms of HSCs.
- ▶ Can be fit to experimental (mouse) data.
- ▶ Some prediction/results about the significance of certain mechanisms.

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- ▶ Model describing central mechanisms of HSCs.
- ▶ Can be fit to experimental (mouse) data.
- ▶ Some prediction/results about the significance of certain mechanisms.
- ▶ A notion of stem cell fitness.

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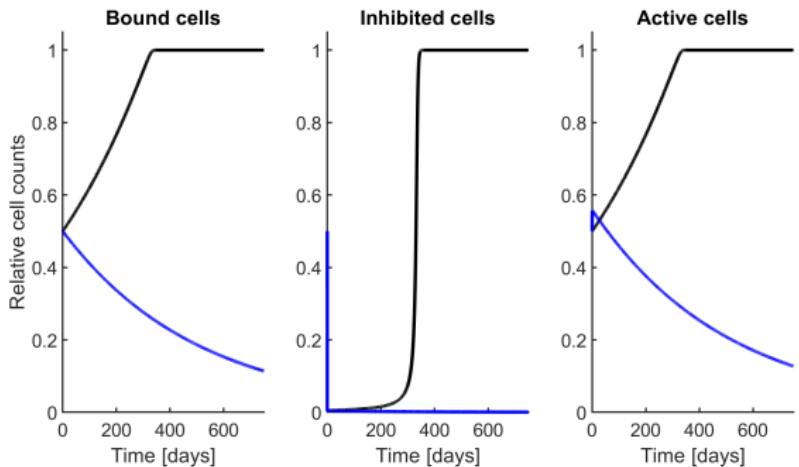
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Reducing the model

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$$\begin{aligned}\frac{dN_E}{dt} &= -b_C N_E I_C + u_C N_C - b_L N_E I_L + u_L N_L \\ \frac{dN_C}{dt} &= b_C N_E I_C - u_C N_C \\ \frac{dI_C}{dt} &= -b_C N_E I_C + 2r_C A_C - d_{I_C} I_C \\ \frac{dA_C}{dt} &= u_C N_C - r_C A_C - d_{A_C} A_C \\ \frac{dN_L}{dt} &= b_L N_E I_L - u_L N_L \\ \frac{dI_L}{dt} &= -b_L N_E I_L + 2r_L A_L - d_{I_L} I_L \\ \frac{dA_L}{dt} &= u_L N_L - r_L A_L - d_{A_L} A_L\end{aligned}$$

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$$\frac{dN_C}{dt} = u_C \left(\frac{2\rho_C(1 - N_C - N_L)}{\alpha_C + 1 - N_C - N_L} - 1 \right) N_C$$
$$\frac{dN_L}{dt} = u_L \left(\frac{2\rho_L(1 - N_L - N_L)}{\alpha_L + 1 - N_L - N_L} - 1 \right) N_L$$

where $\alpha_C = \frac{d_{I_C}}{b_C K}$, $\alpha_L = \frac{d_{I_L}}{b_L K}$, $\rho_C = \frac{r_C}{r_C + d_{A_C}}$ and $\rho_L = \frac{r_L}{r_L + d_{A_L}}$

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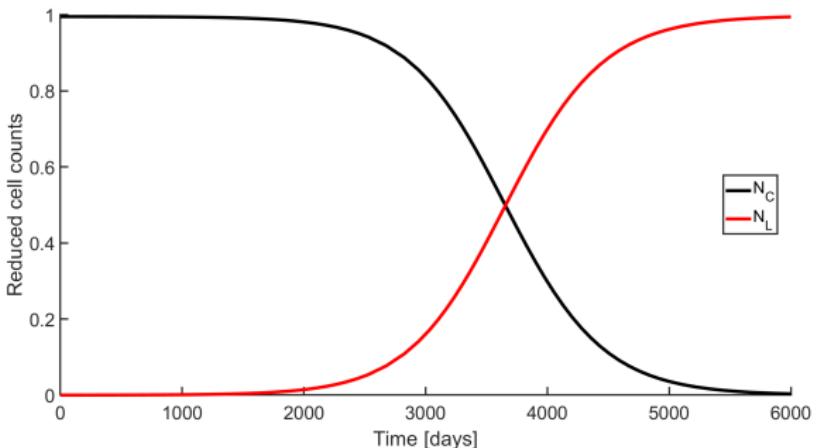
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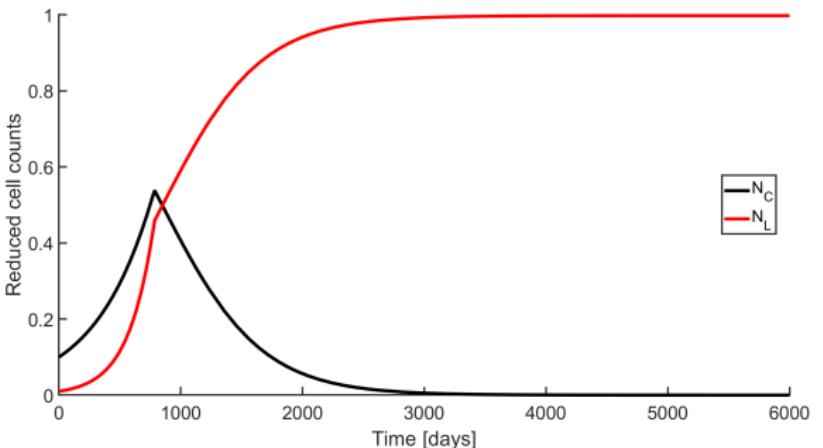
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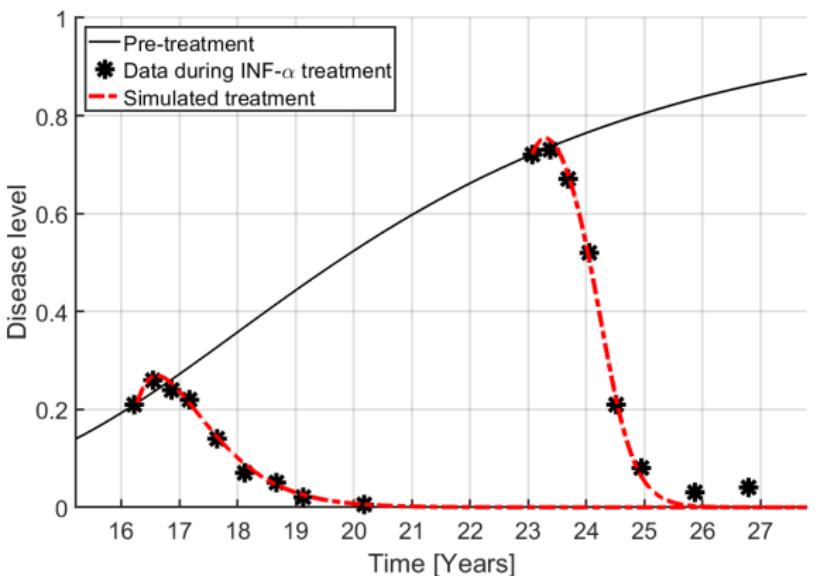
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- Models allow us to test hypotheses.

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- ▶ Models allow us to test hypotheses.
- ▶ Mathematical modelling of stem cells help shine a light on a system which is otherwise hard to investigate.

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- ▶ Models allow us to test hypotheses.
- ▶ Mathematical modelling of stem cells help shine a light on a system which is otherwise hard to investigate.
- ▶ In particular: Limited self-renewal with “recharging” through the niche leads to certain properties of the HSC-bone-marrow system, and a notion of HSC fitness.

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- ▶ HSC fitness is cell intrinsic, i.e. it could be possible to determine it from studies involving only single cell types.

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- ▶ HSC fitness is cell intrinsic, i.e. it could be possible to determine it from studies involving only single cell types.
- ▶ By considering feedback signalling from blood, fitting with and perhaps even predicting patient-data could be possible.

Thank you for your attention.

Any questions?



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Bonus figure

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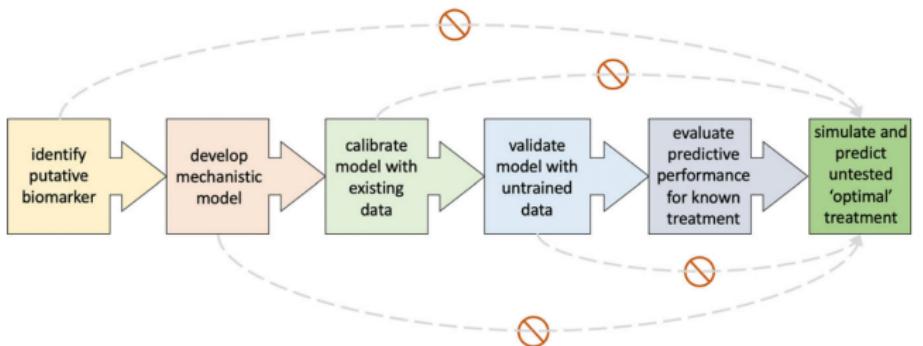
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(Brady and Enderling, 2019)