



University of Milan Department of Physics

Overshoot of cancer stem cell population

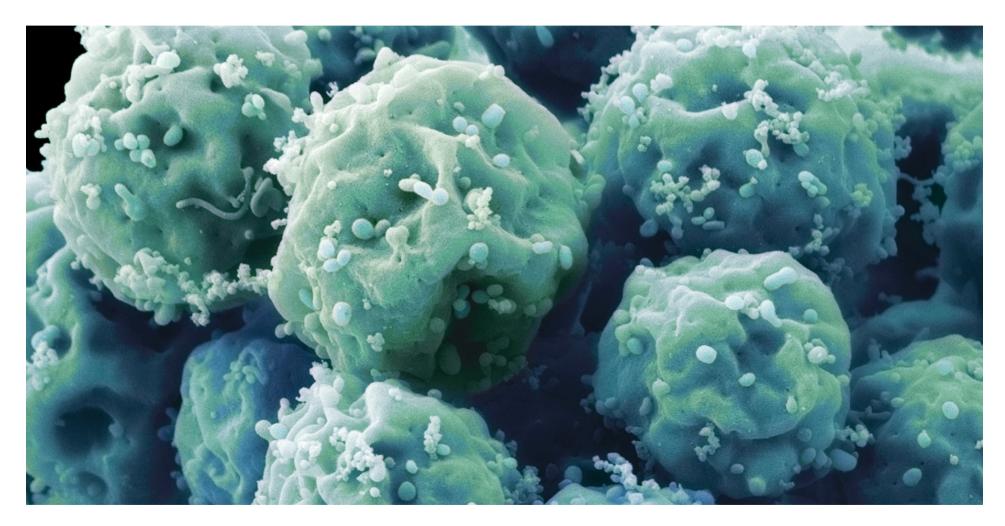
A theoretical model via rate eq. and stochastic simulations

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Unusually for physicists, the main protagonist of our talk shall be biological cells... in particular CANCER STEM CELLS (CSC)

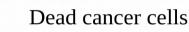


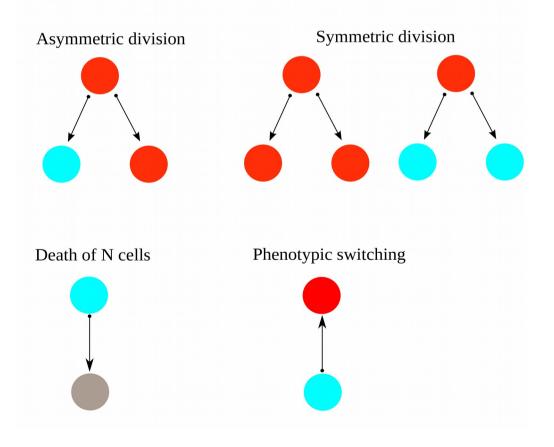
The Bayer Scientific Magazine

Aberrant CSC Hierarchy Theory for Tumor Growth

Cancer Stem Cells (CSC)

Normal cancer cells (N)



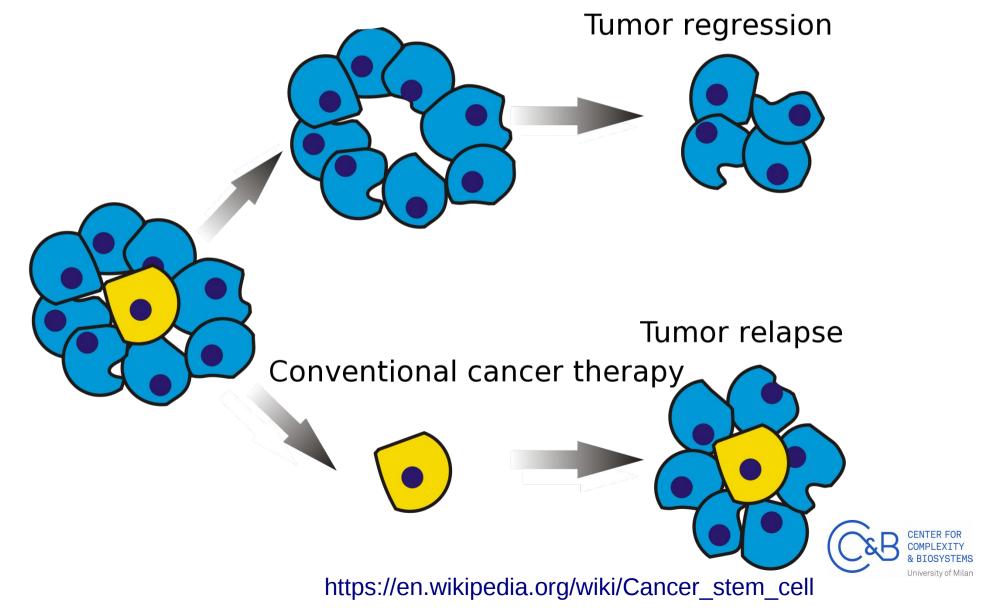


- only small fraction of malignant cells drives tumor growth (stem cell like);
- CSC prop.: self-renewal, differentiation into multiple cell types, longer life span;
- origin not clear;
- first identified in leukemia, then also in brain, breast, colon, pancreas cancers and melanoma.



The hierarchy model suggests a specific therapy to eradicate CSC population

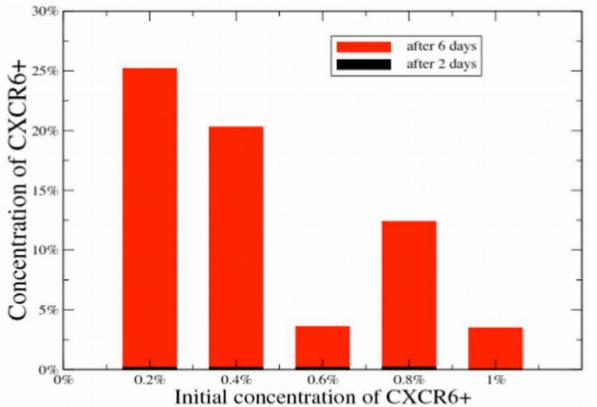
Cancer stem cells specific therapy



Howewer, experiments performed by La Porta's Group in the Department of Biosciences suggest this may not always be the case



Observed Overshoot of CSC Population in a Melanoma (initially reduced under 1% of total)

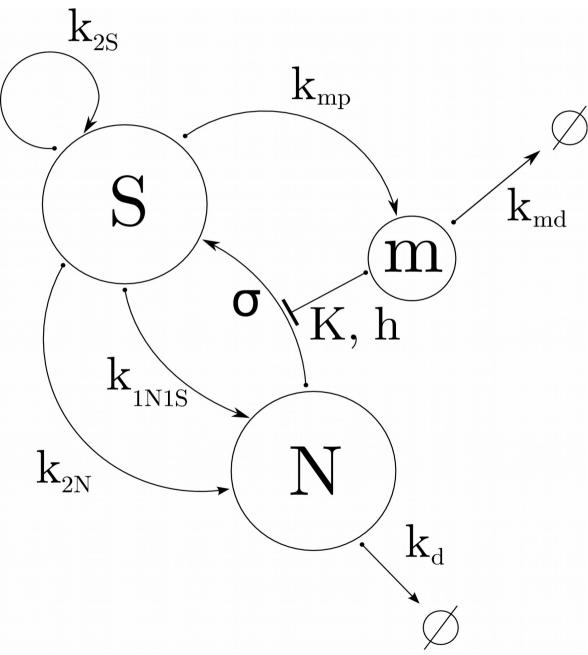


The fraction of CSC grows from less than 1% of total cells to a maximum of 25% after some days and then decreases to a steady value.

Sellerio et al., Sci. Rep., 2015



Reaction Network of the model





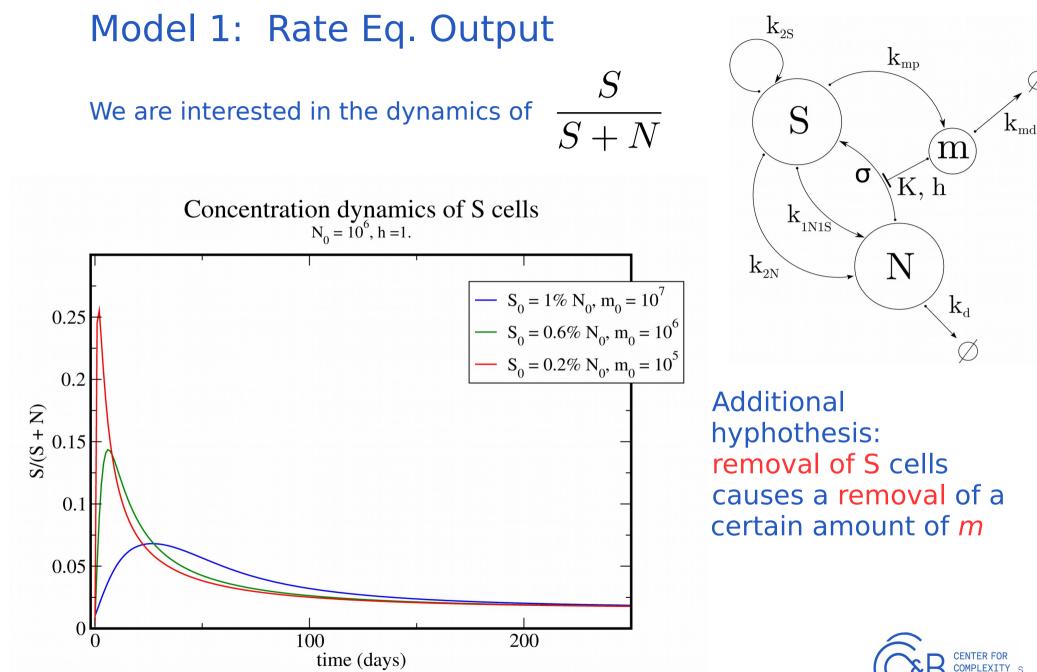
Model 1: Rate Equations

Initial Values: $m(0) = m_0^{-1}$, $S(0) = S_0^{-1}$, $N(0) = N_0^{-1}$.

$$\begin{cases} \dot{m} = k_{np}S - k_{nm}m \\ \dot{S} = (k_{2S} - k_{2N})S + \sigma N \left(1 - \frac{m^{h}}{k^{h} + m^{h}}\right) \\ \dot{N} = (2k_{2N} + k_{1N1S})S - \left(k_{d} + \sigma \left(1 - \frac{m^{h}}{k^{h} + m^{h}}\right)\right)N \end{cases}$$

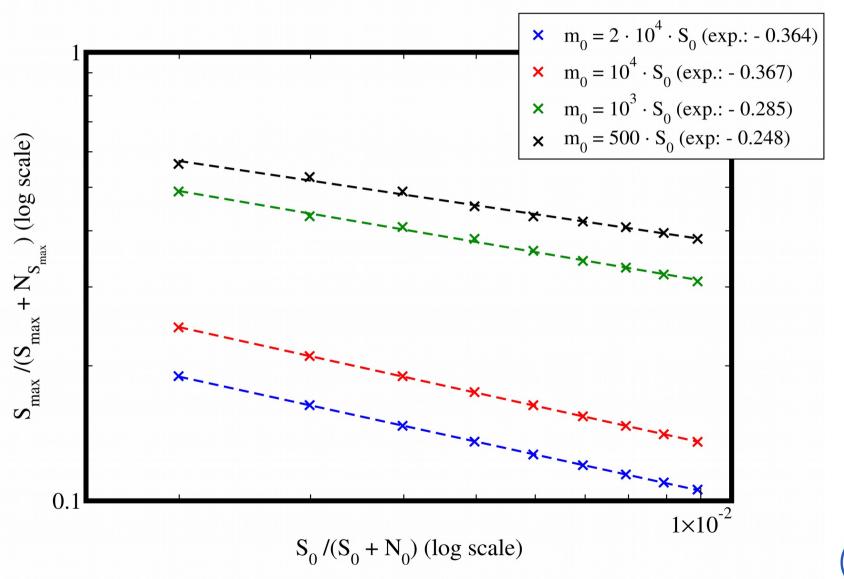
Simple model;presence of many unknown parameters.





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Power law for maxima of concentration of S cells with respect to initial concentration (for different ratios S_0/m_0)





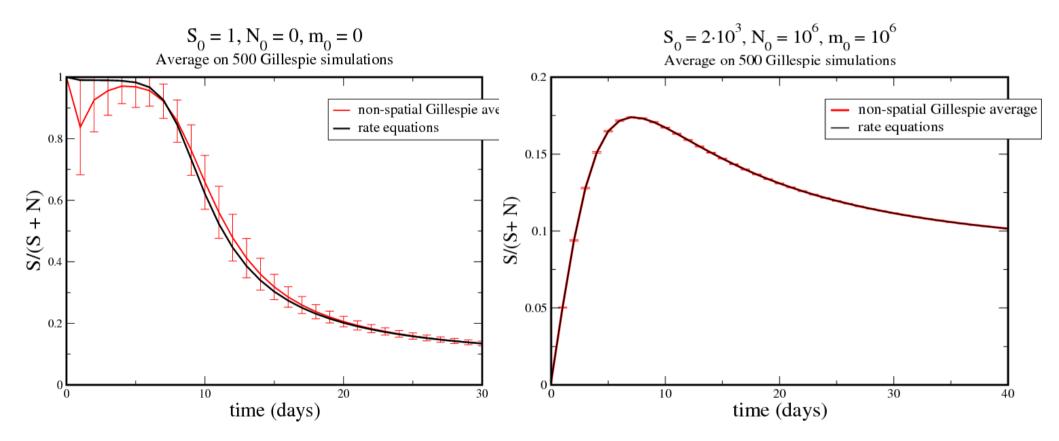
To improve our model, consider as a first step:

- the discrete nature of cells and molecules;
- possibility of reactions with low number of cells and molecules involved (failure of well-mixed system hyphothesis in most biochemical systems);
- stochastic noise in biochemical reactions.

This lead to modeling the proposed reaction network via the non-spatial Gillespie algorithm (computational consistent solution of the chemical master equations).



Model 2: Gillespie average vs rate equations

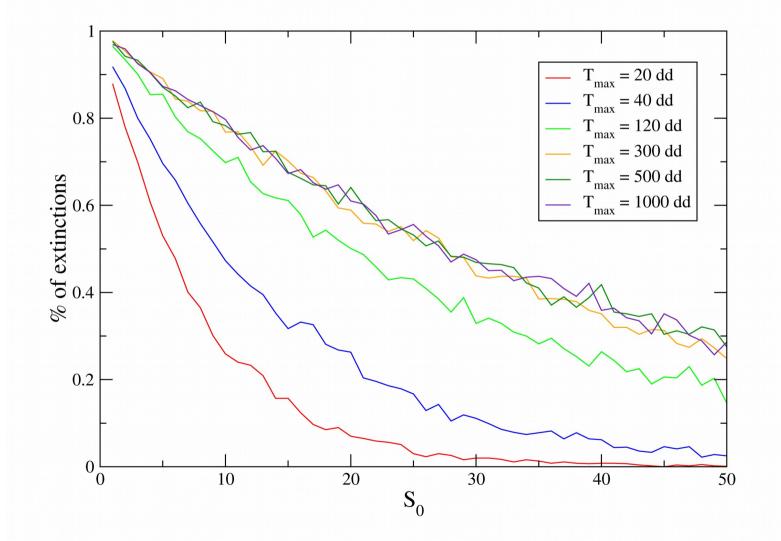


For high rates of spontaneous extinction of S cells dynamics, averages shall not match rate equations (intrinsic stochastic nature of the system).



Spontaneous extinction of tumor dynamics in absence of switch (small $\varepsilon = k_{2S} - k_{2N}$)

% of spontaneous extinctions in absence of switching ($\sigma = 0$)





Next step: modeling in space

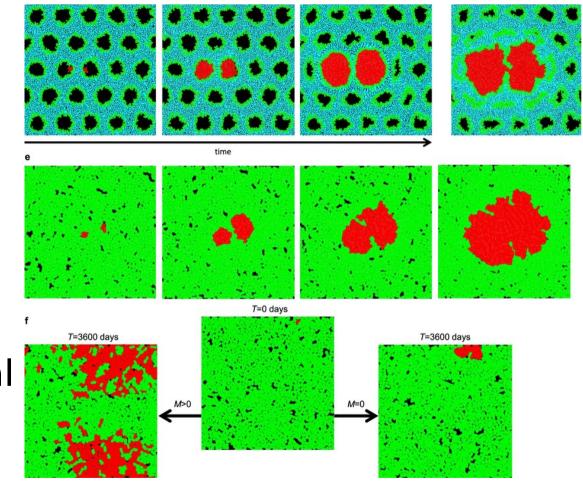
- Phenotypic switch dependent on local cell--molecule interactions;
- molecule m are produced in S cells sites and then diffuse;
- Effects of crowded environment on cell division rates.



Future Developments: Spatial Gillespie (2d and 3d)

Role of diffusion of molecule *m*;

- Stochastic fluctuations in space;
- Detection of spatial patterns to match images observed at the microscope.



Waclaw et al., Nature, 2015.

