

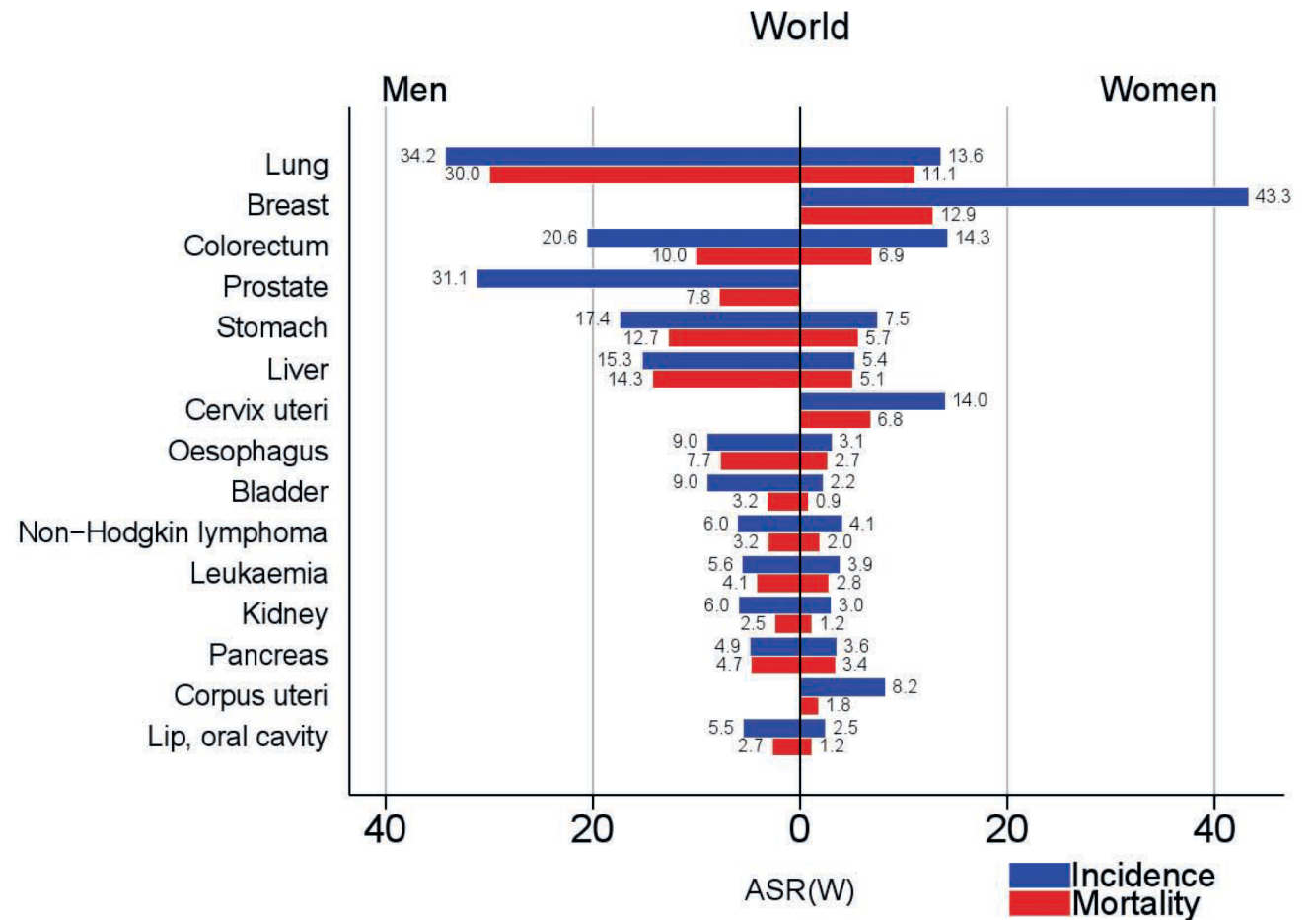
Logical modelling and analysis of cell adhesion properties along Epithelial to Mesenchymal Transition

Gianluca Selvaggio

Workshop on Logical Modelling of Cellular Networks – ECCB18

Intro

Cancer is a leading cause of death worldwide with **8.8M** deaths in 2015.



ASR incidence and mortality per 100 000, by major sites, in men and women, 2012.

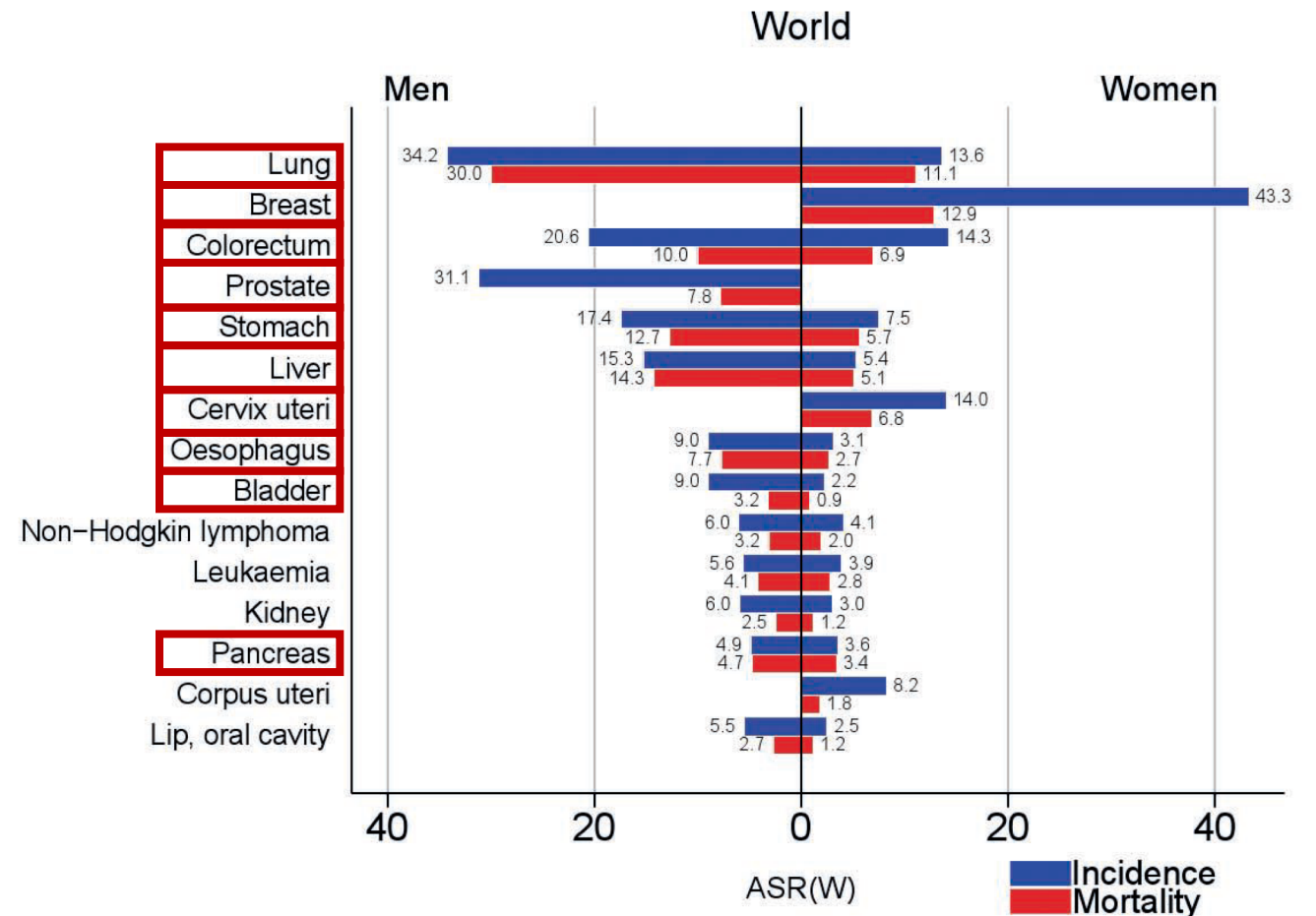
GLOBOCAN 2012 v1.0, Cancer Incidence and Mortality Worldwide: IARC Cancer Base No. 11

<http://www.who.int/en/news-room/fact-sheets/detail/cancer>

Intro

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Carcinoma: cancers that arise from **epithelial** cells.



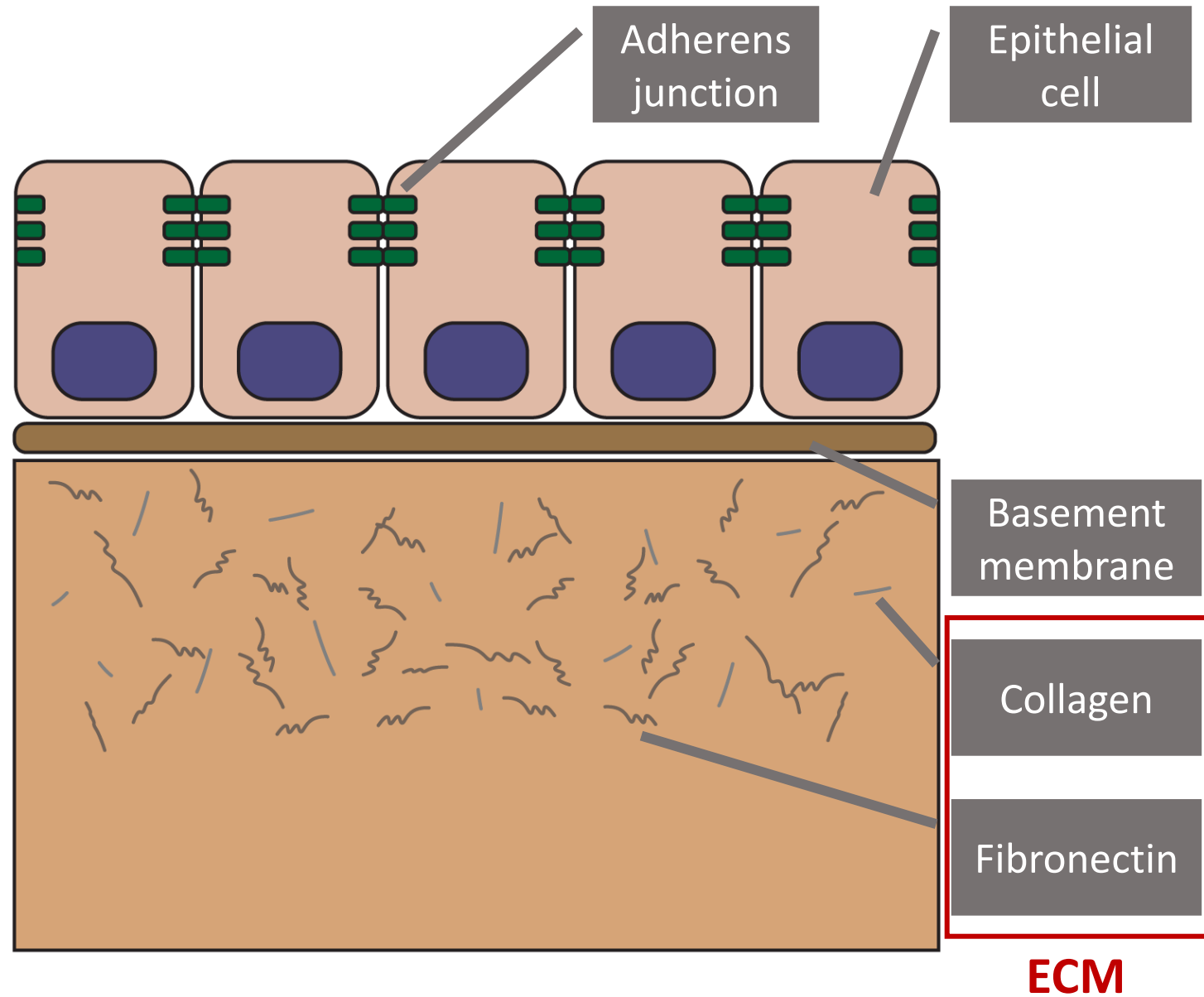
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Epithelium

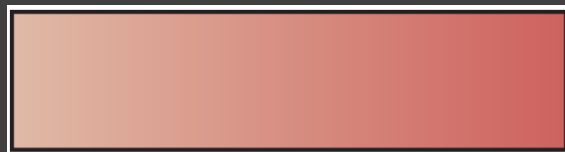
Thin layers of cells that cover internal/external surfaces of bodies and organs.



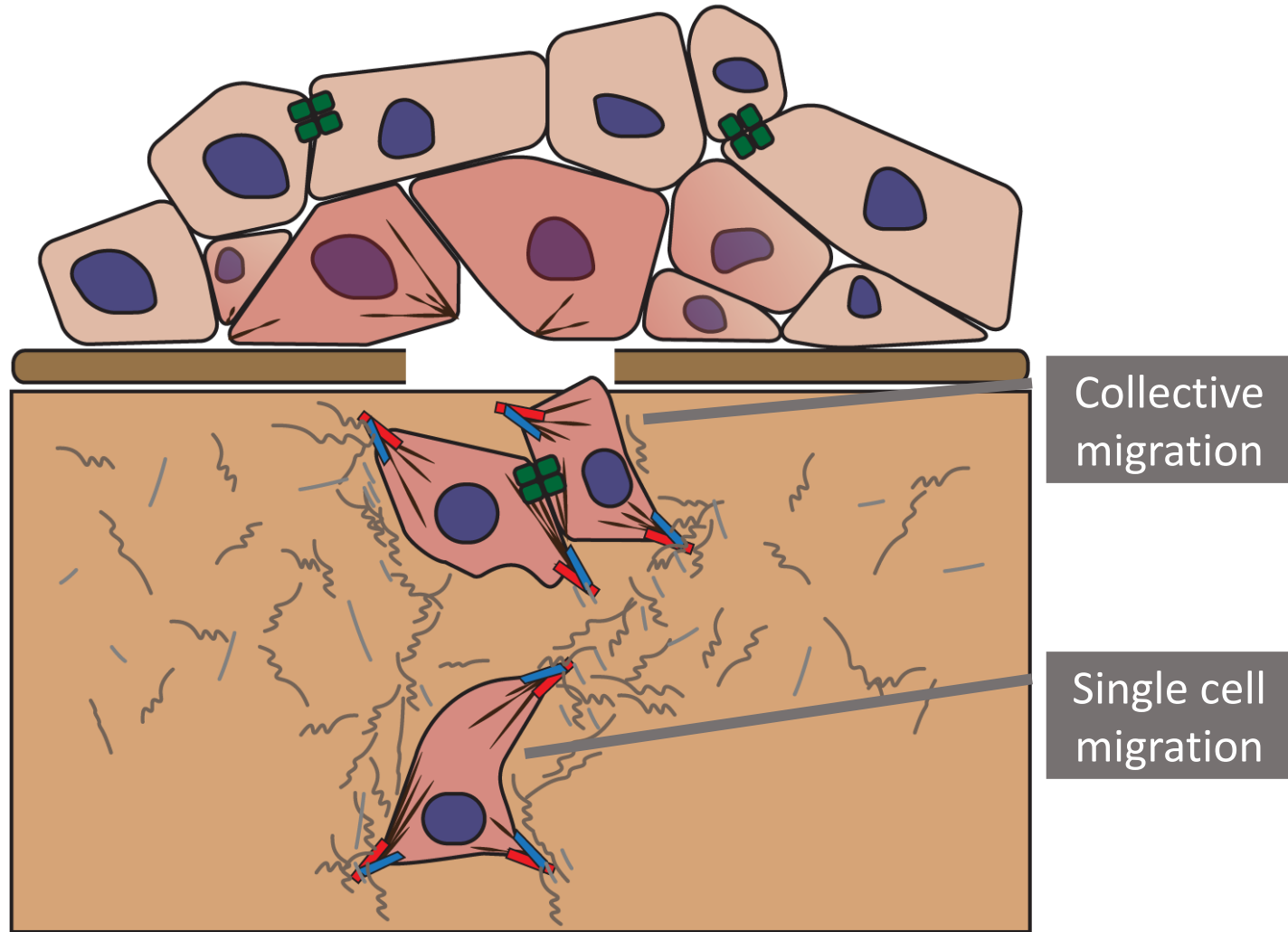
EMT: invasion and metastasis

Bidirectional signalling between cancer cells and the tumour microenvironment drives the progressive loss of **epithelial** properties combined with the cumulative acquisition of **mesenchymal** features (**EMT**).

Epi



Mes



We propose a logical modelling approach to **investigate** and **understand** the mechanism at play during **EMT**, and the influence of the **tumour environment** on **cell adhesion properties**.

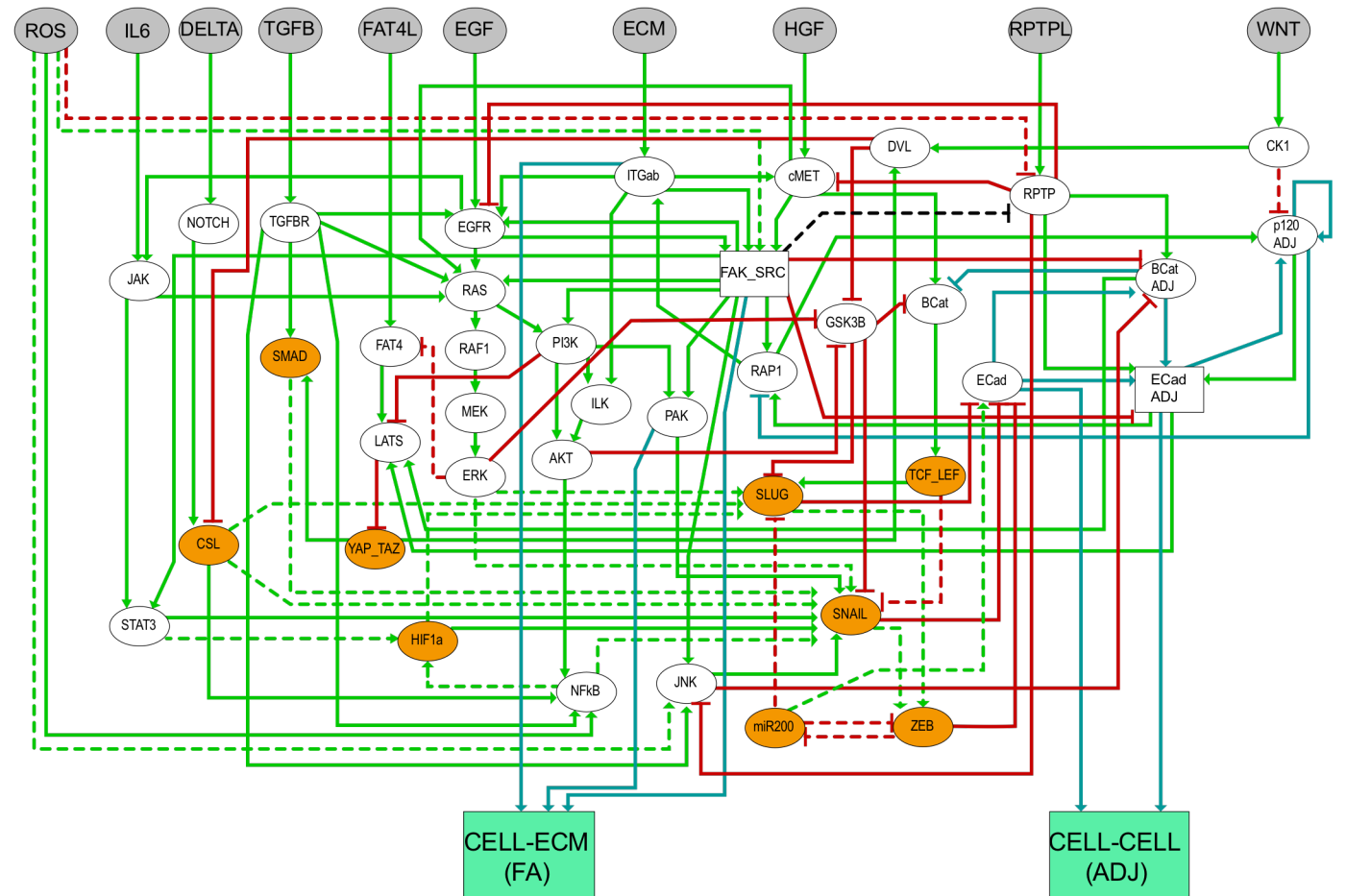
Logical formalism

The complexity and dimension (**components**) of the molecular network combined with a **lack of quantitative** information on **kinetic parameters, concentrations** and **mechanistic** insights on protein interactions motivate the use of logical modelling.

Boolean/Multivalued abstraction: each regulatory component is associated to a discrete variable representing its levels of activity, of concentration, etc. → **functional level.**

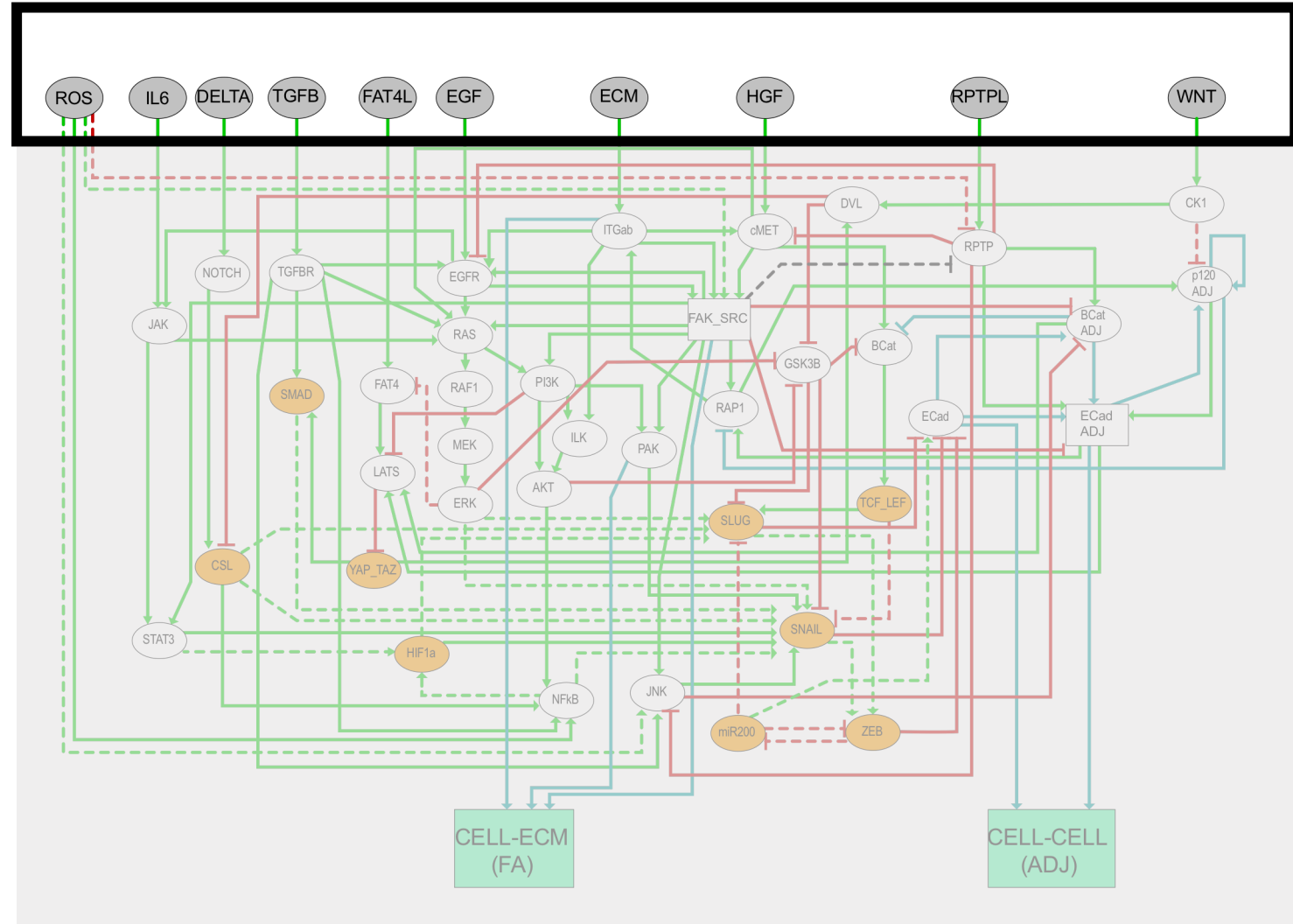
Each **regulatory component** is associated to a set of incoming interactions defining the evolution of the corresponding variable.

Model of cell adhesion properties



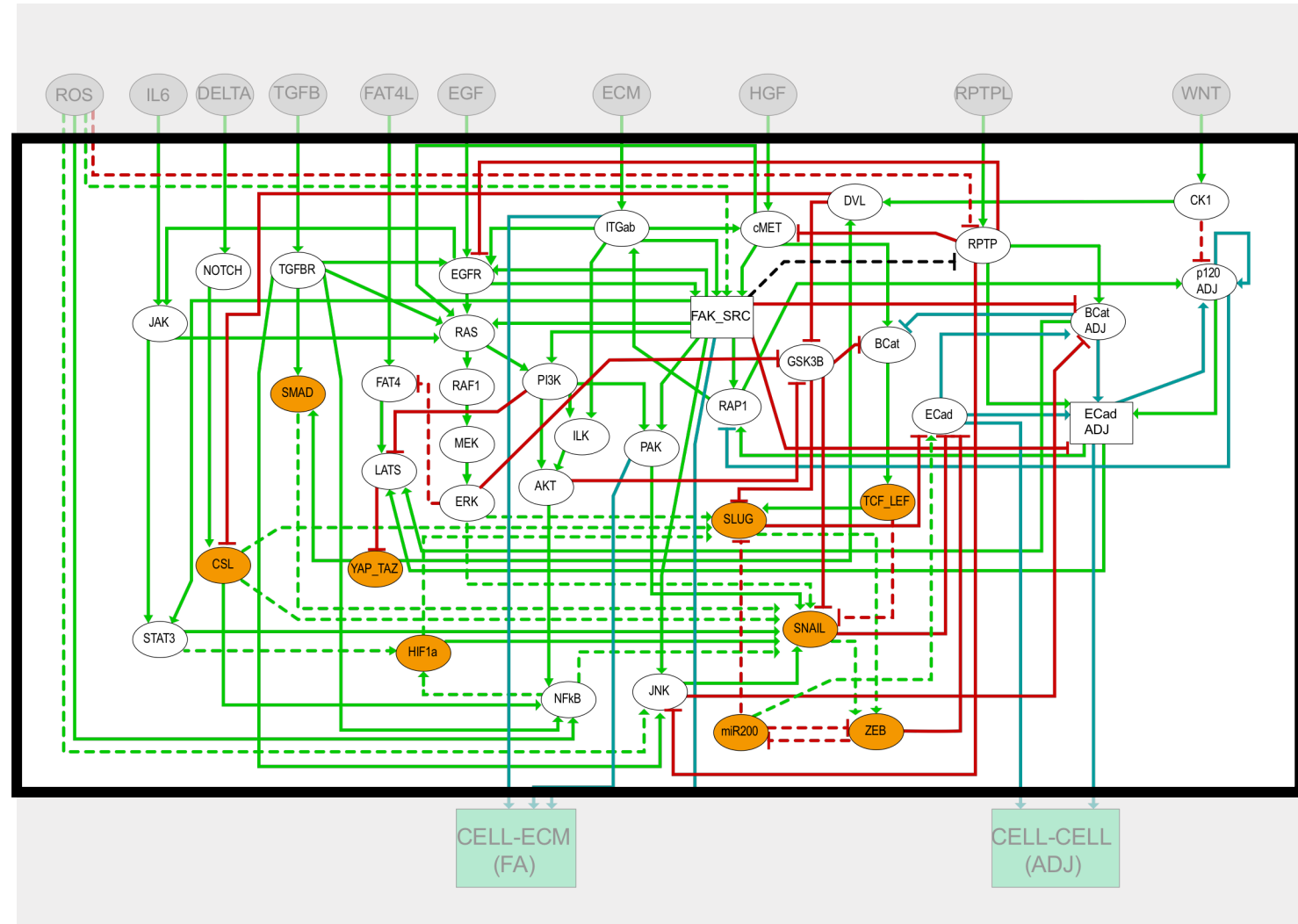
Model of cell adhesion properties

Input (e.g. growth factors, cell contacts, cytokines etc.)



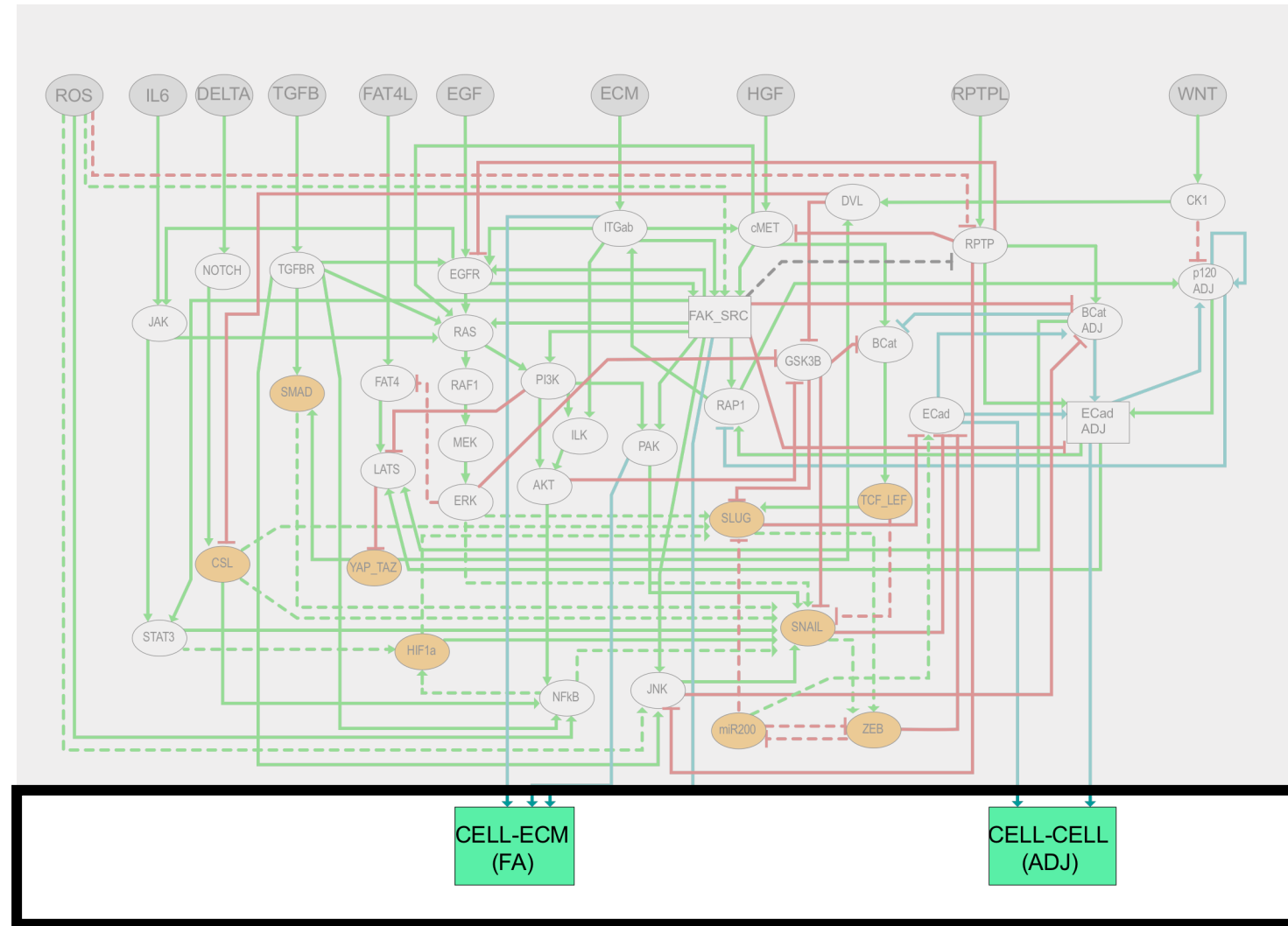
Model of cell adhesion properties

Internal components (e.g. kinases, transcription factors etc.)

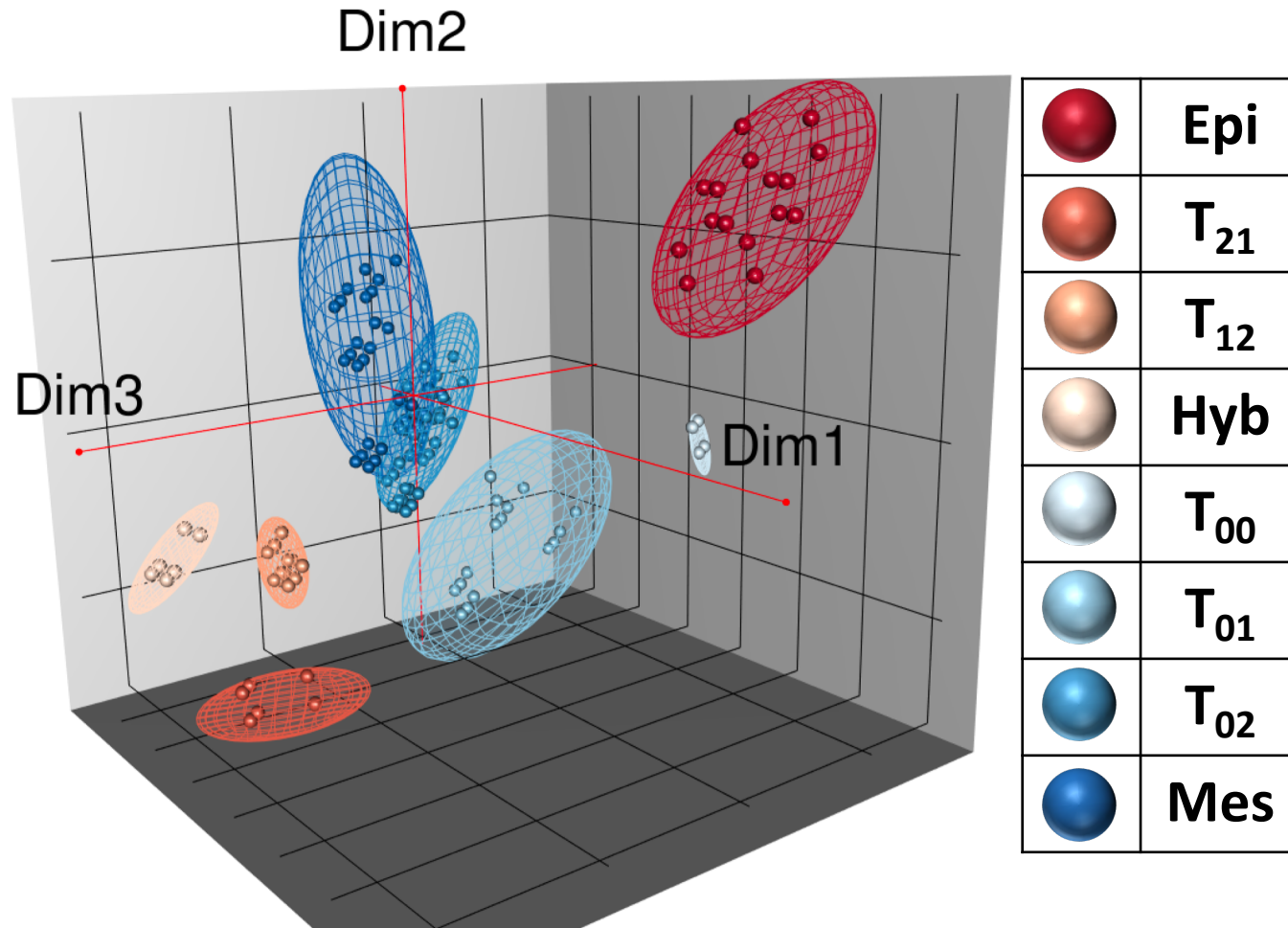


Model of cell adhesion properties

Outputs



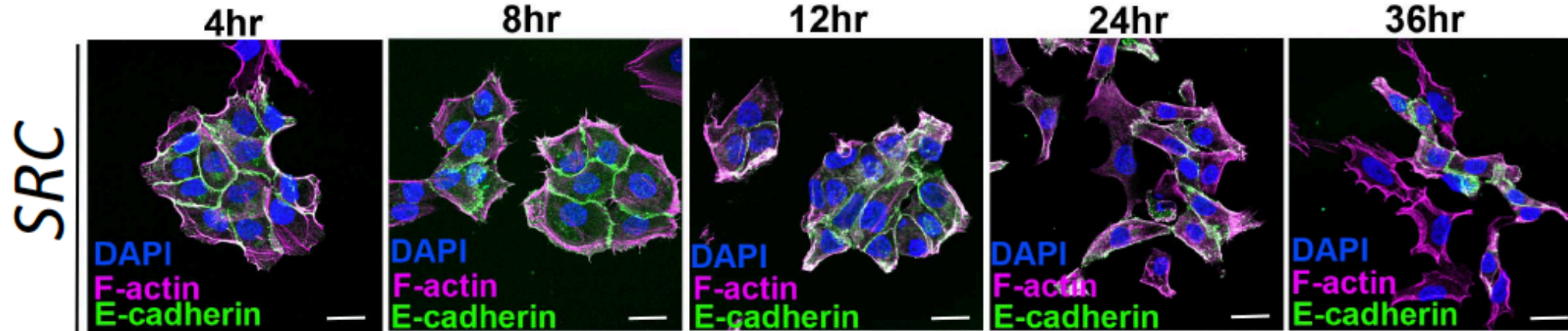
Asymptotic behaviours



Is there a clear difference between the hypothetical phenotypes?

Model Predictions

SRC is a proto-oncogene tyrosine kinase whose activation is capable of transforming **non-tumorigenic** epithelial breast cell line MCF10A.



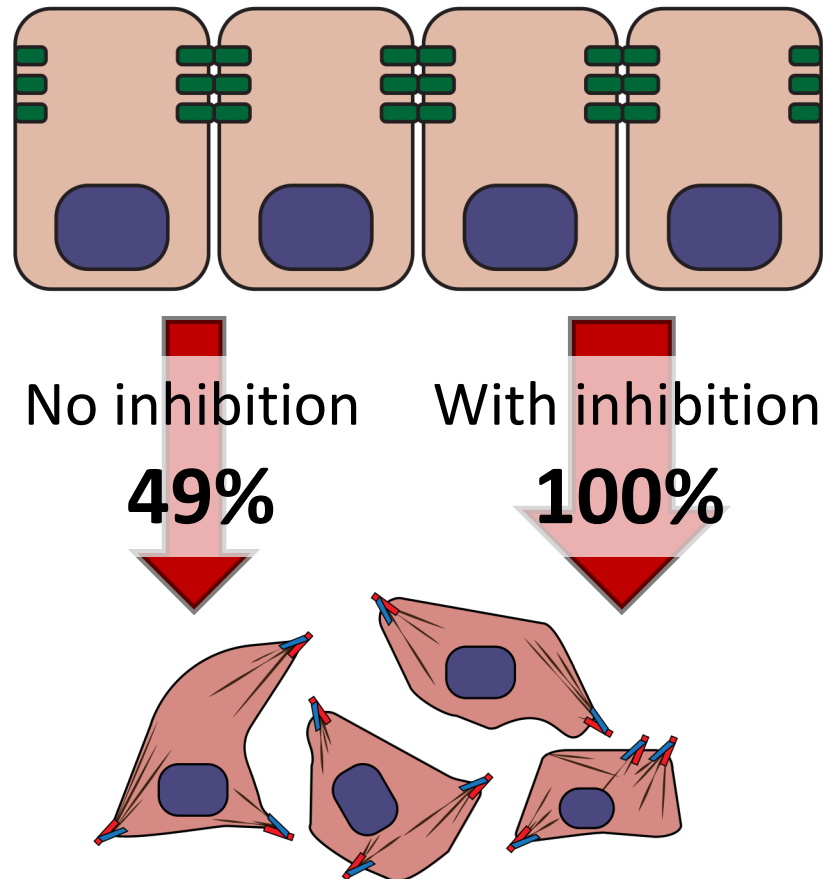
With the first version of the model only ~**49%** of the simulations starting from an **epithelial** state reached the **mesenchymal** phenotype

Model Predictions

μ-array data from literature (Hirsch, H.A. *et. al.* 2010) suggested **SRC inhibition of PTPR** (cell contact activated phosphatase).

Model Predictions

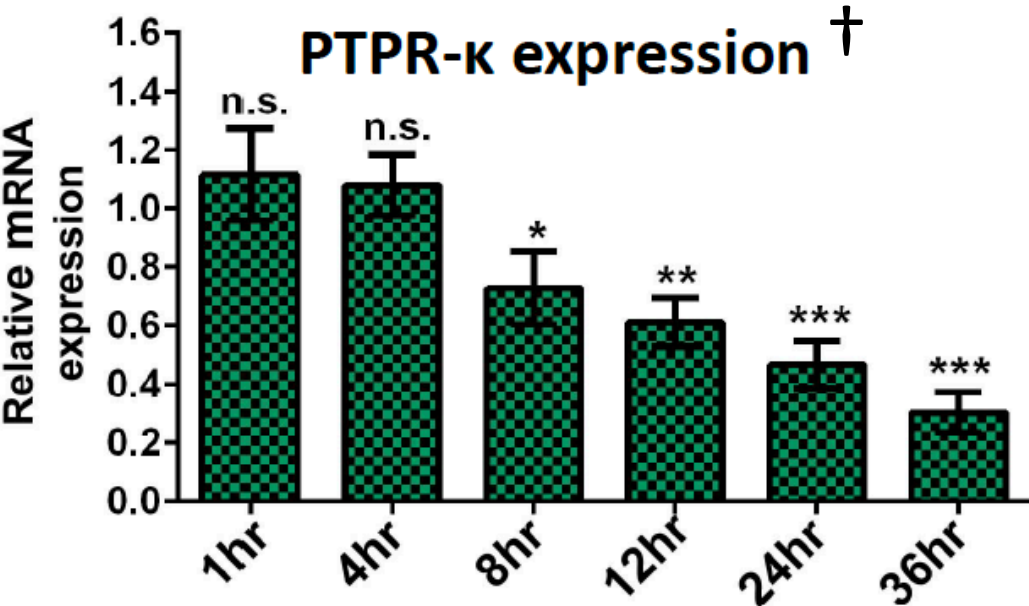
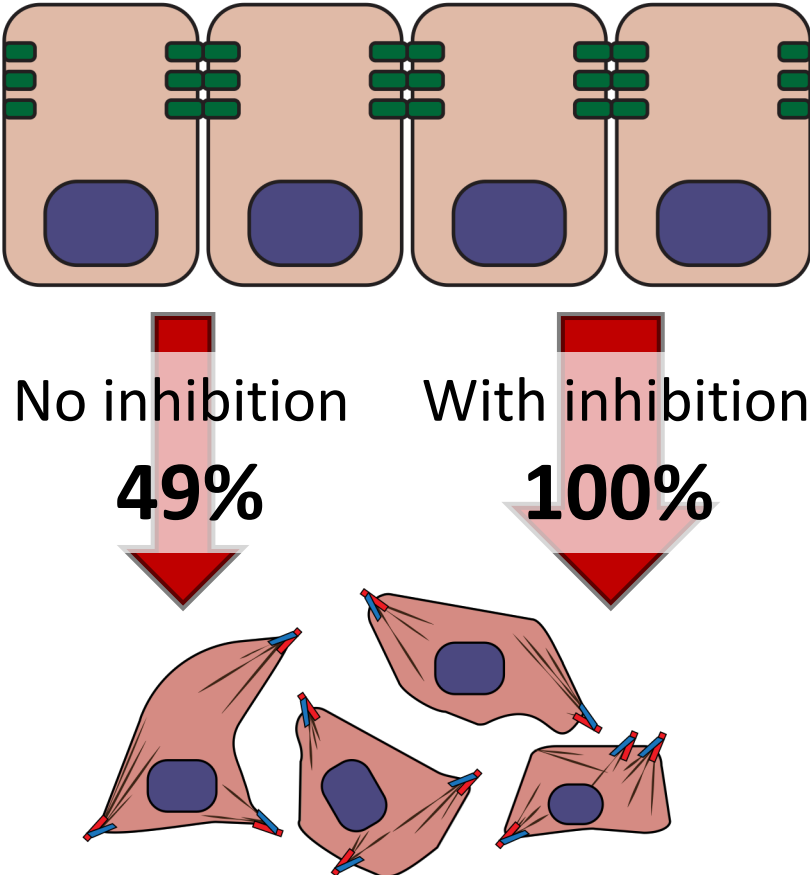
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Hirsch, H.A. *et. al.* – Cancer Cell. 17(4) - (2010)

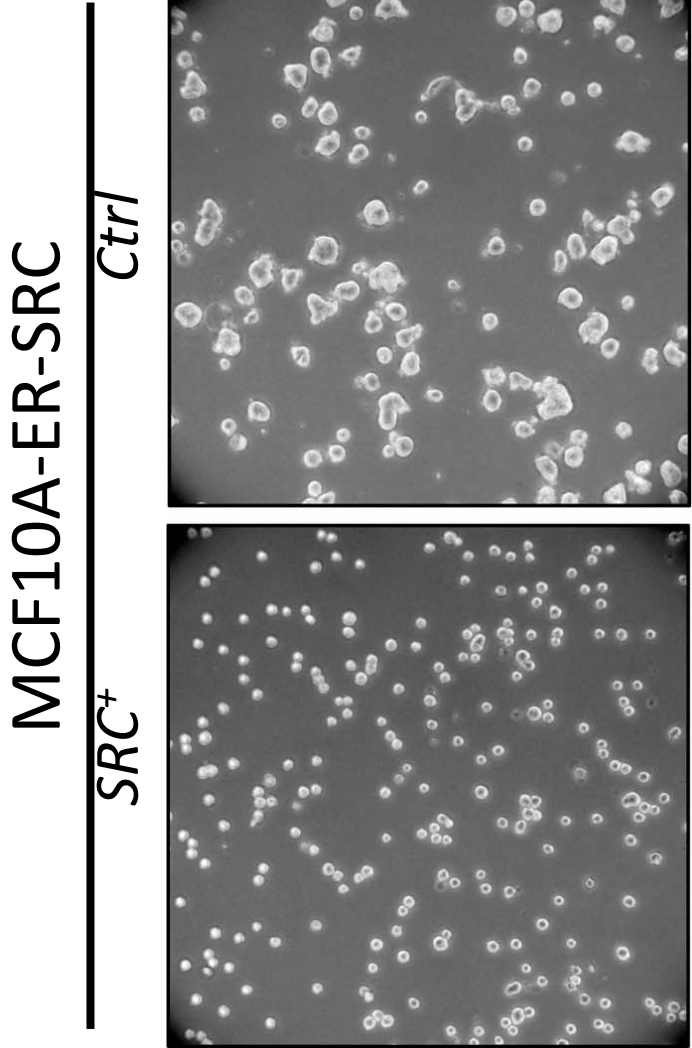
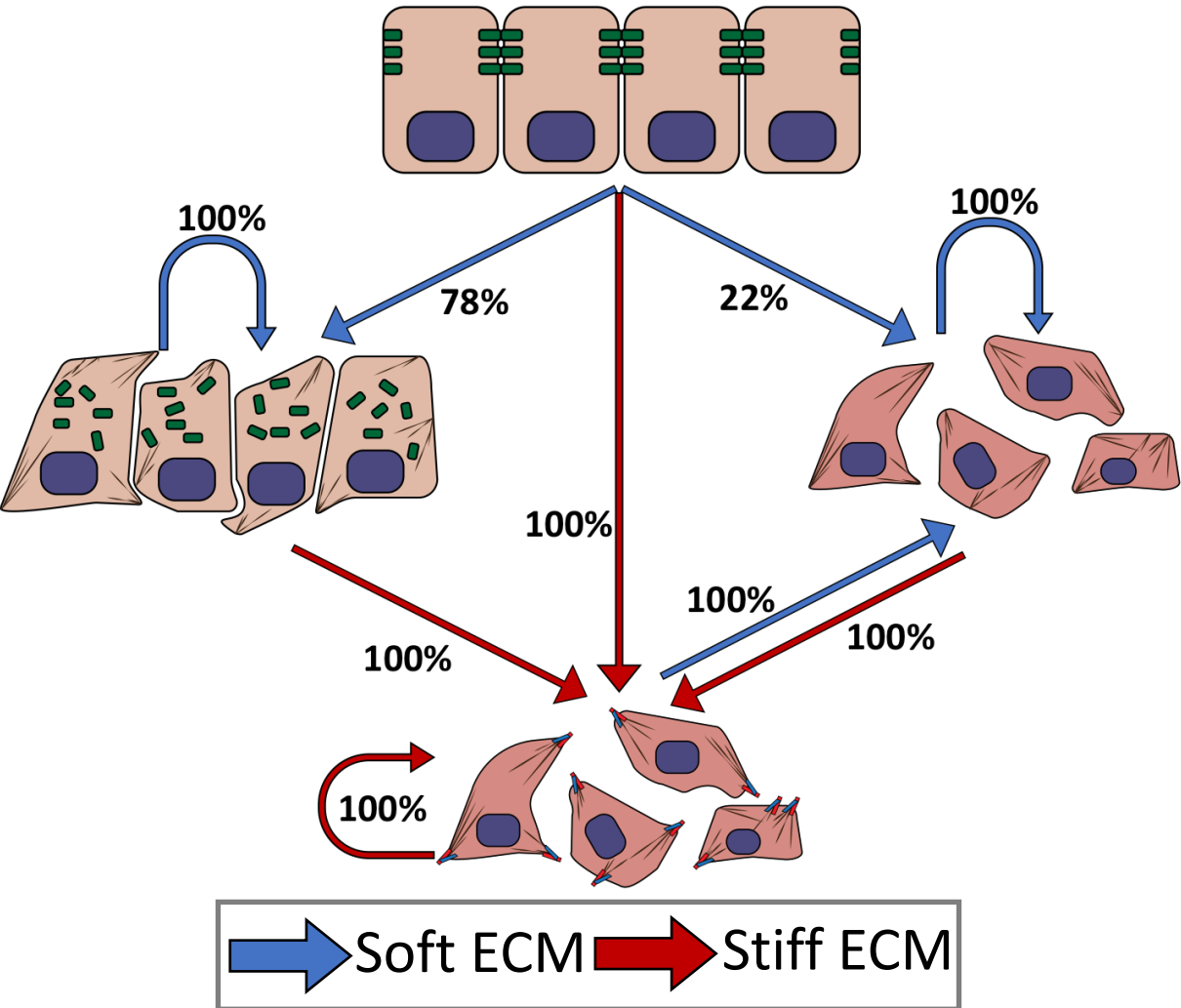
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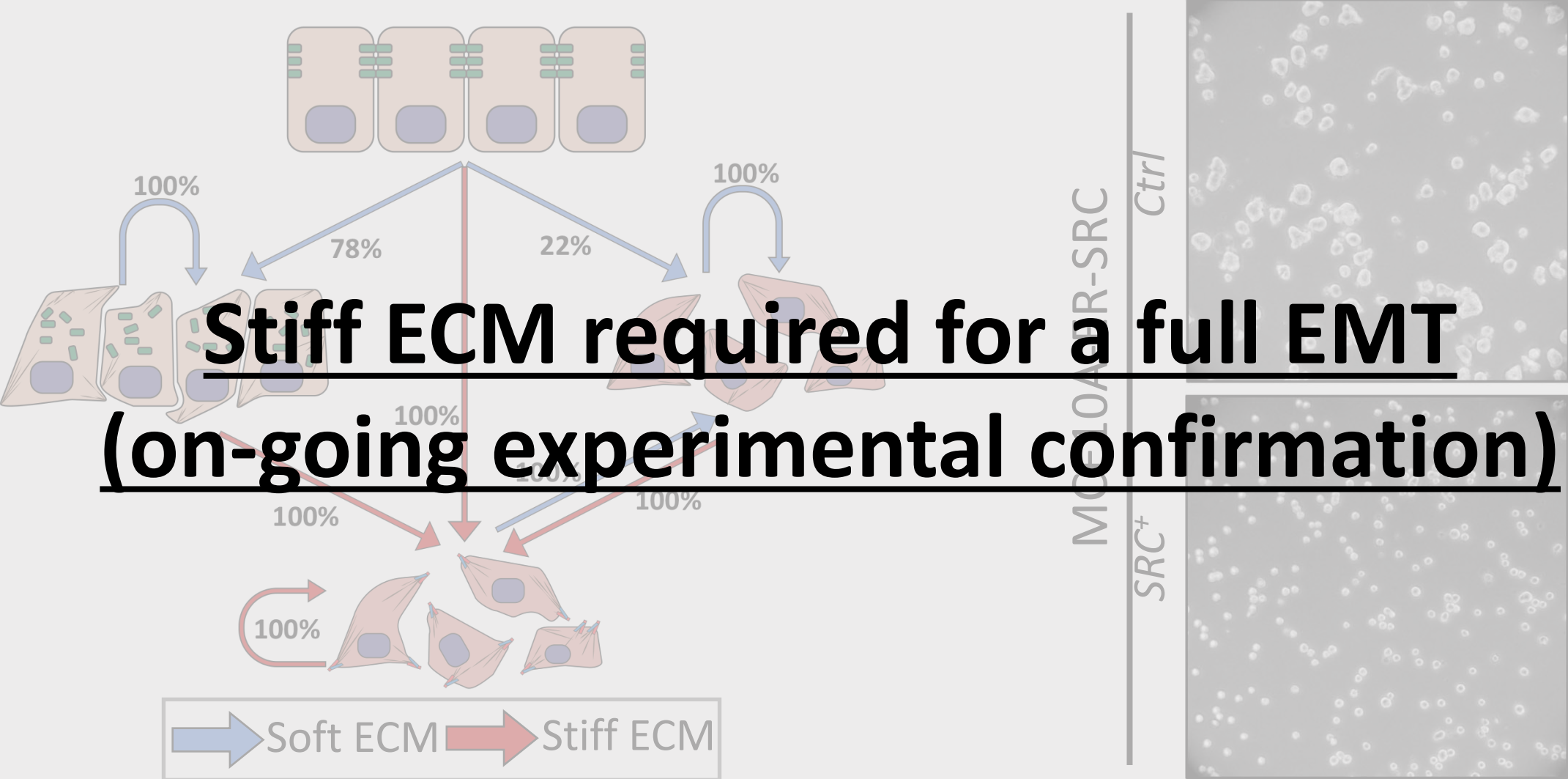


Hirsch, H.A. *et. al.* – Cancer Cell. 17(4) - (2010)
[†] In-house validation performed by A. Pawar

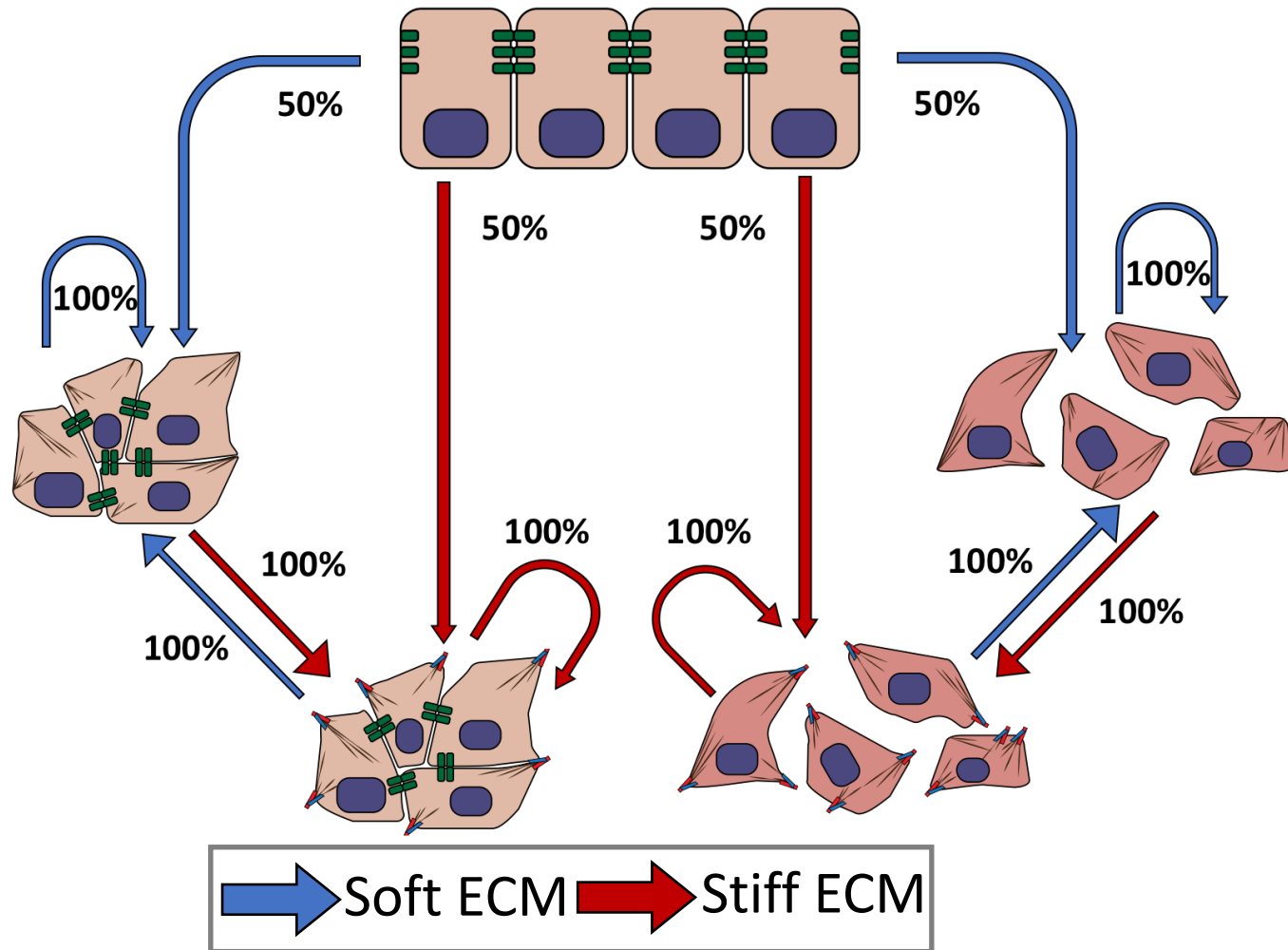
Model Predictions: *In silico* vs *in vitro*: SRC⁺



Model Predictions: *In silico* vs *in vitro*: SRC⁺

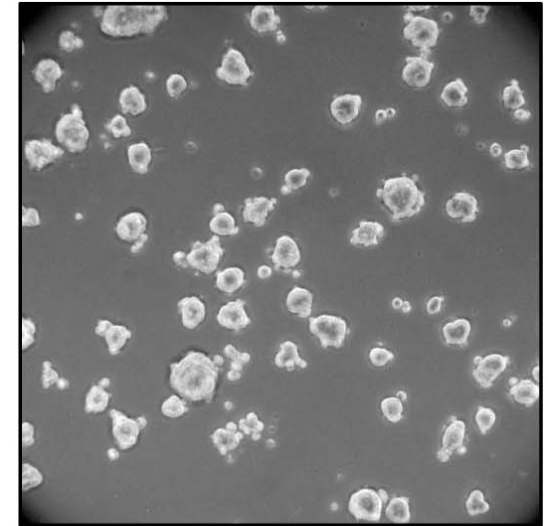


Model Predictions: *In silico* vs *in vitro* $SRC^+ / PTPR^+$

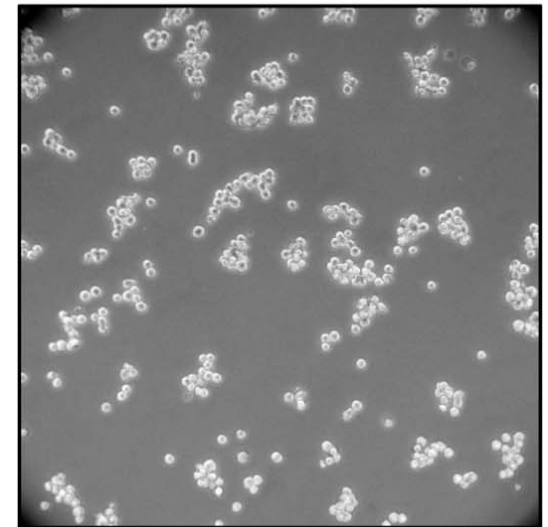


MCF10A-ER-SRC|PTPR⁺

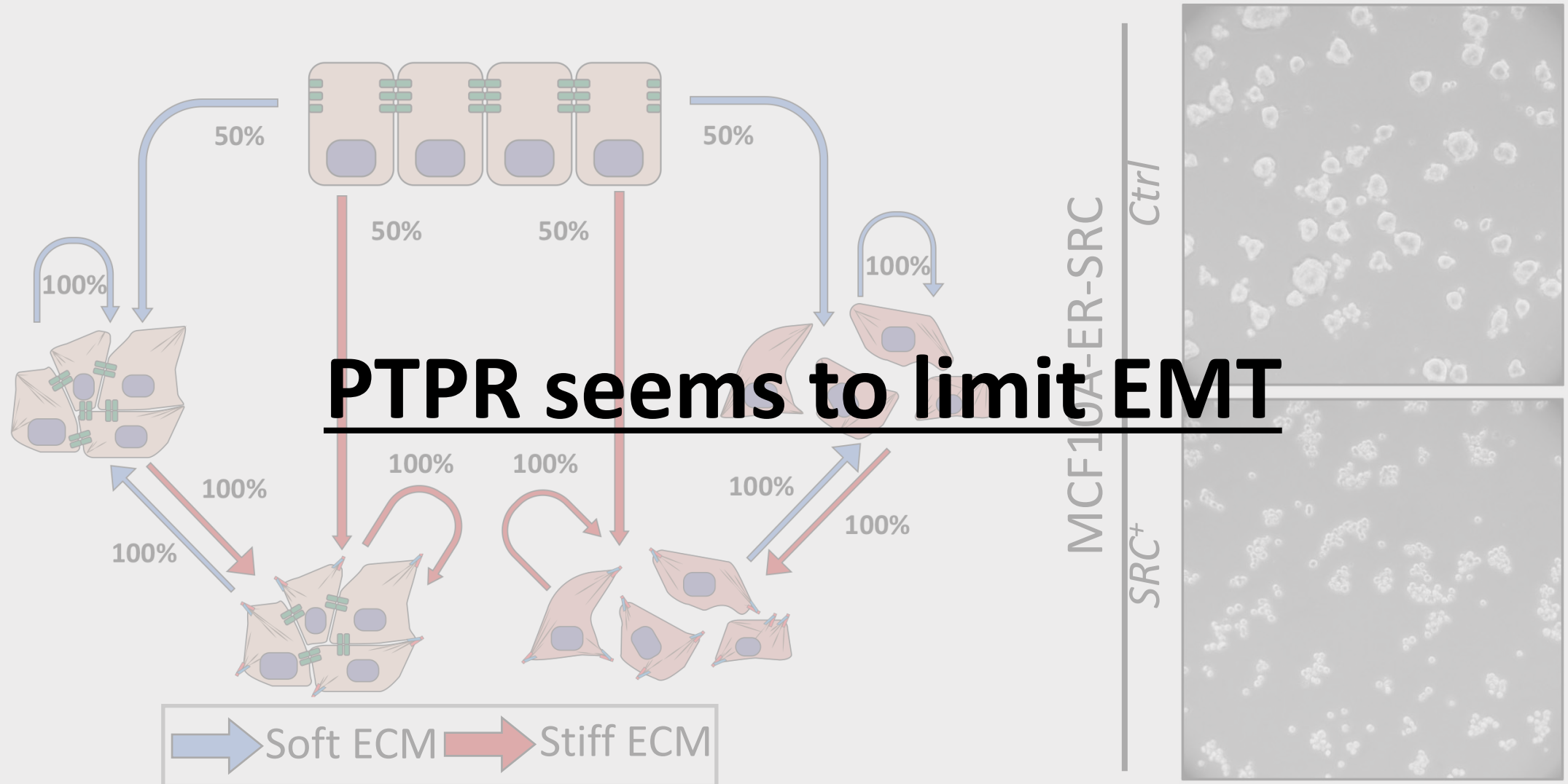
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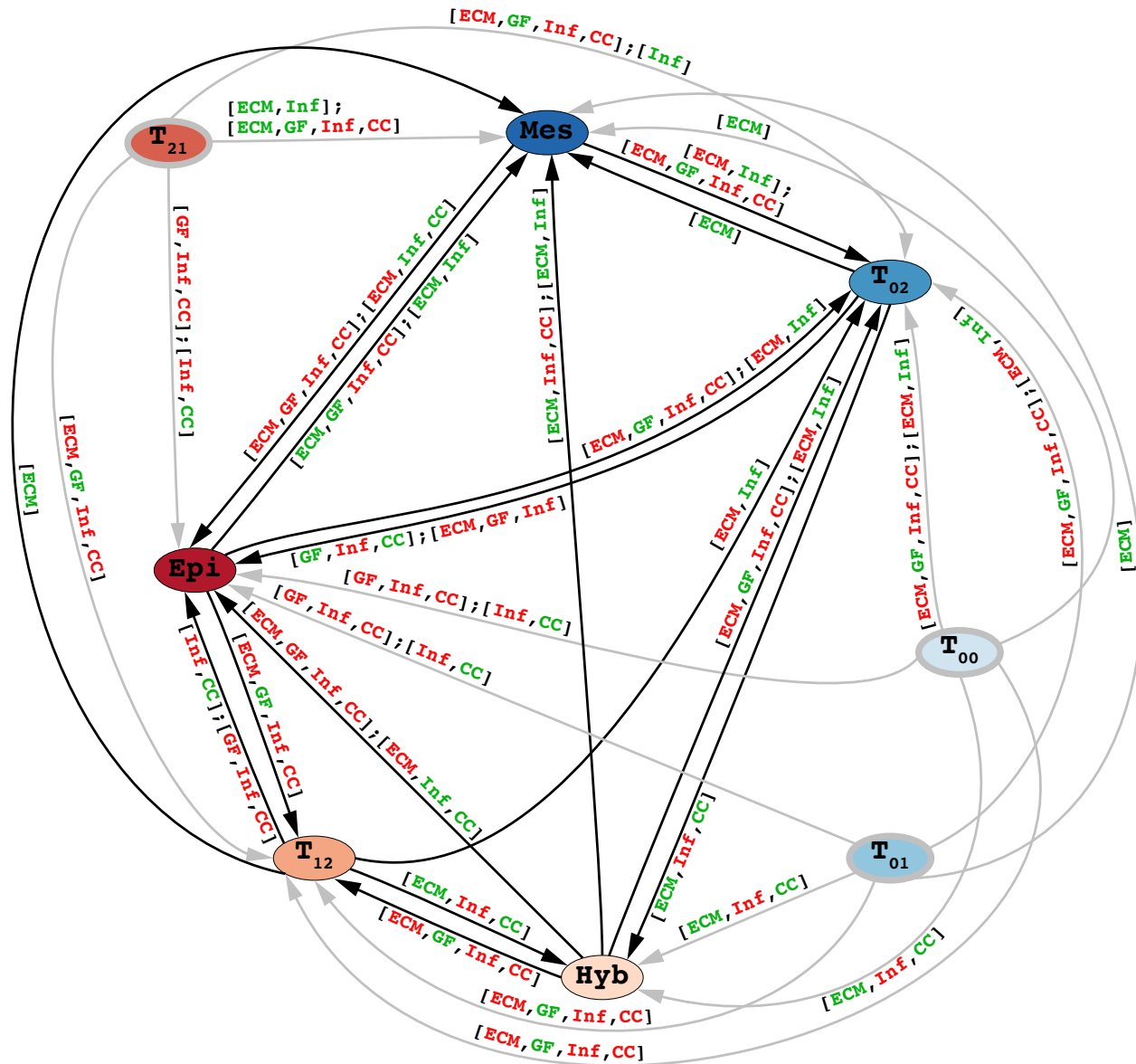
SRC⁺



Model Predictions: *In silico* vs *in vitro* $SRC^+ / PTPR^+$



We used model-checking techniques to assess environmental influence:



Model Predictions: Phenotype Plasticity

ECM	Growth Factors (GF)	Inflammation (Inf)	Cell-Cell contact (CC)
ECM	EGF, HGF	IL6, ROS, TGFβ	RPTPL, FAT4L

Conclusions

- We provide a **tool** for **probing** in silico **cellular responses** to internal and environmental perturbations
- **PTPR** might be a critical **EMT inhibitor** downstream of *SRC*, by limiting the mesenchymal phenotype and favouring the emergence of hybrid phenotype

Future prospects

- Model **extension** to investigate the link between **EMT** and acquisition of **stemness** features
- Embedding the model in a **multi-cellular** context to unravel interplay between **neighbouring cells**

Acknowledgments

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