

Network dynamics driving cancer metastasis: from design principles to therapeutic approaches

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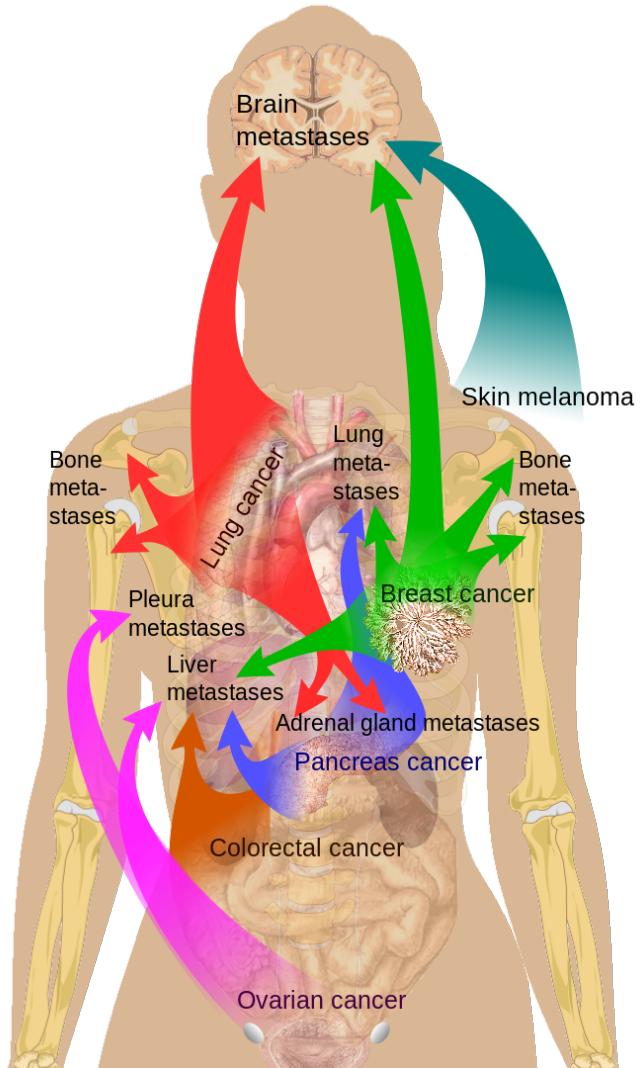
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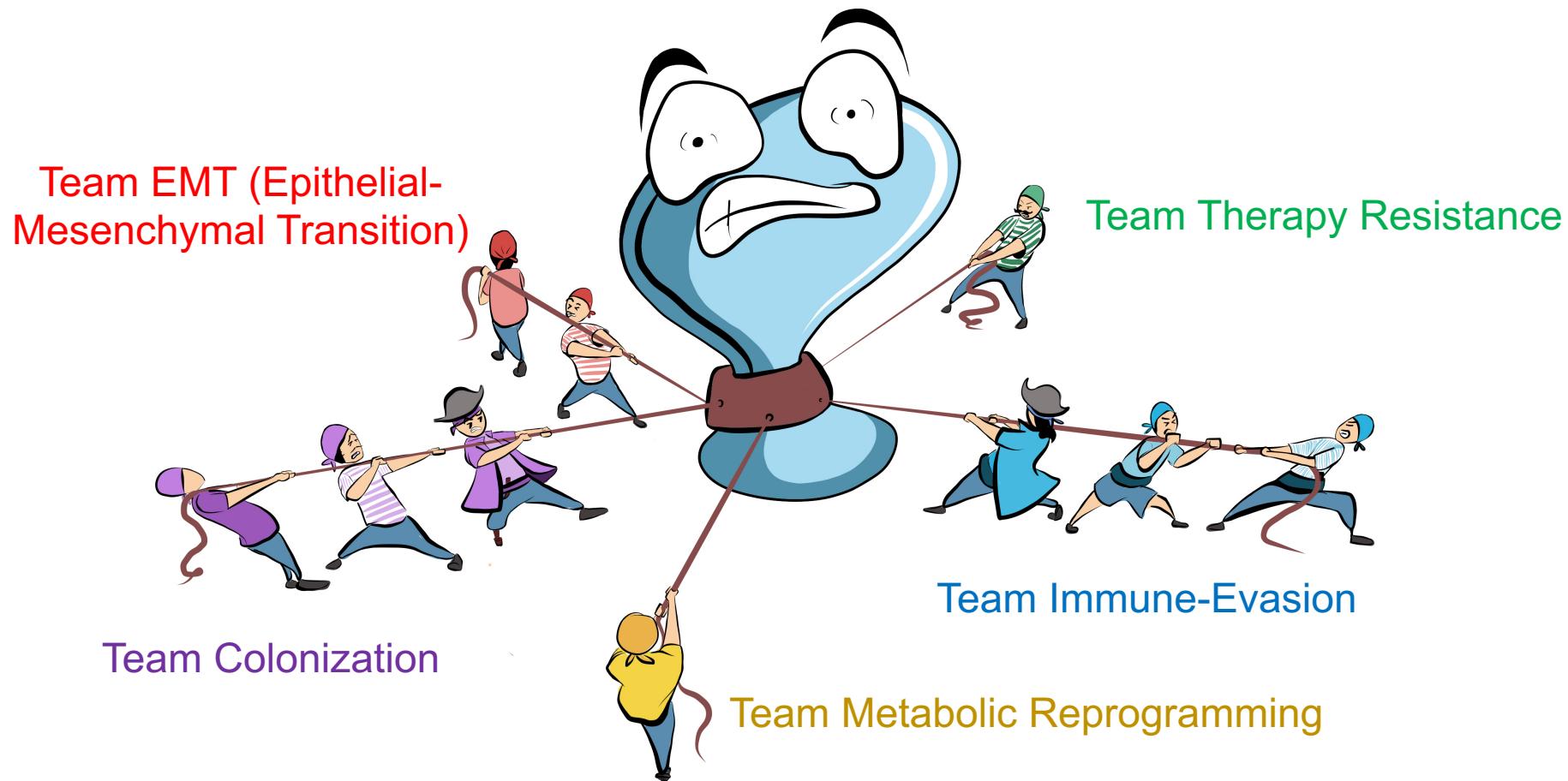
Metastasis : the cause of 90% of all cancer deaths



Metastasis has extremely high attrition ($> 99.9\%$) rates.



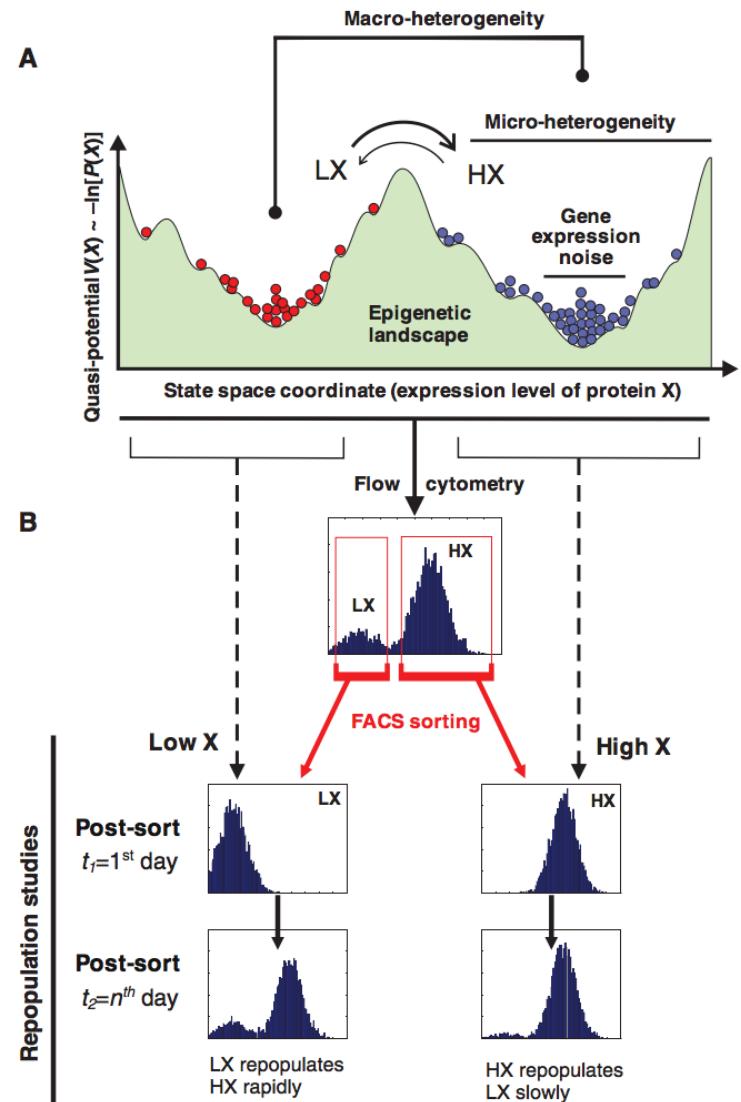
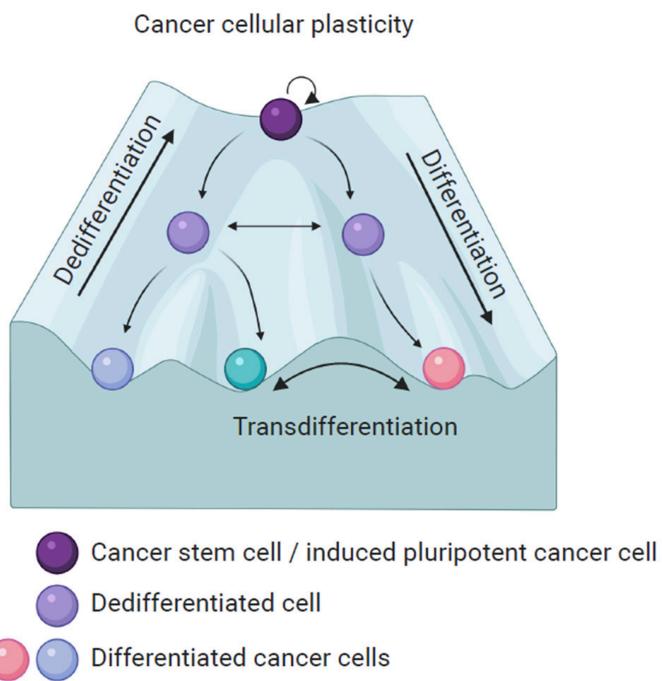
What traits cells need to successfully metastasize?



We need a **dynamic and systems-level understanding** of the process to identify how cells alter these multiple traits together

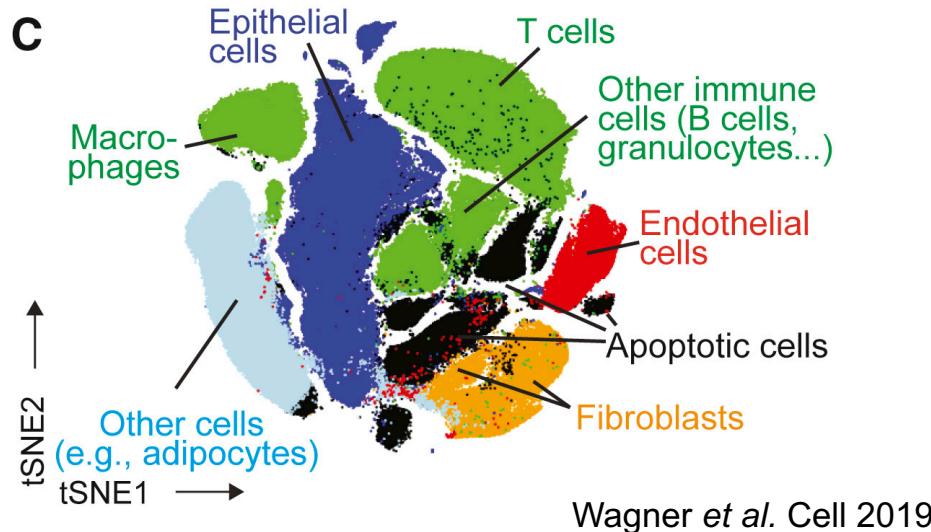
Phenotypic plasticity and non-genetic heterogeneity

Cellular/Phenotypic plasticity:
Ability of cells to switch their
phenotype/behavior
reversibly in response to
environmental conditions

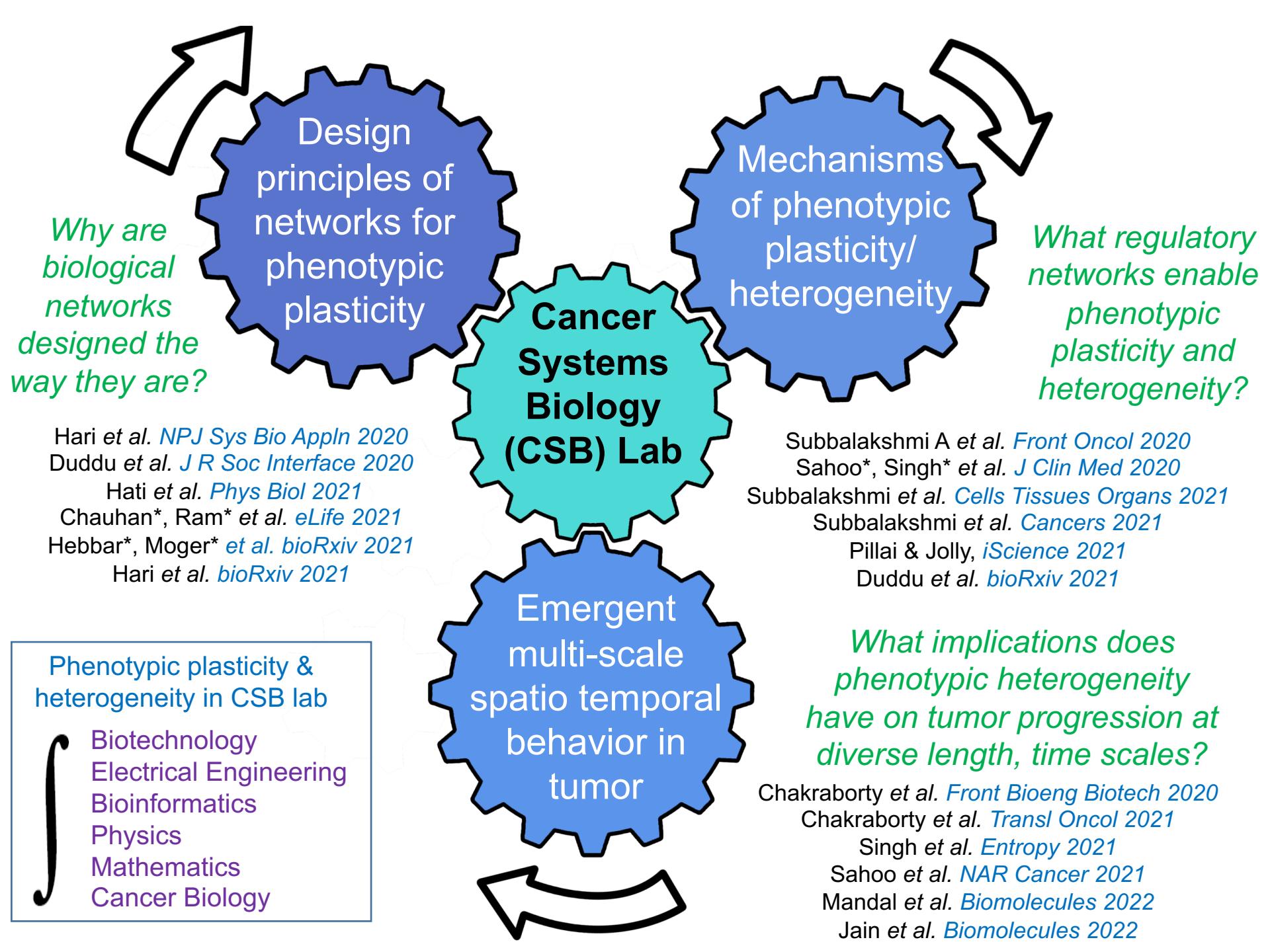


Huang et al. Development 2009
Granados et al. Int J Mol Sci 2020

Open questions about plasticity & heterogeneity in cancer



- How many states can cancer cells exist in?
- How do they switch among these states?
- How do they coordinate behavior among these different axes?
- Can we suggest ways to control plasticity and heterogeneity in a dynamic evolving system?





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Adithya Chedere
Saurav Kumar



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Andrew Armstrong



TIFR Centre for
Interdisciplinary
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Basil Thurakkal



Rik Thompson
Sugandha Bhatia



Pritha Ray
Ajit Dhadve

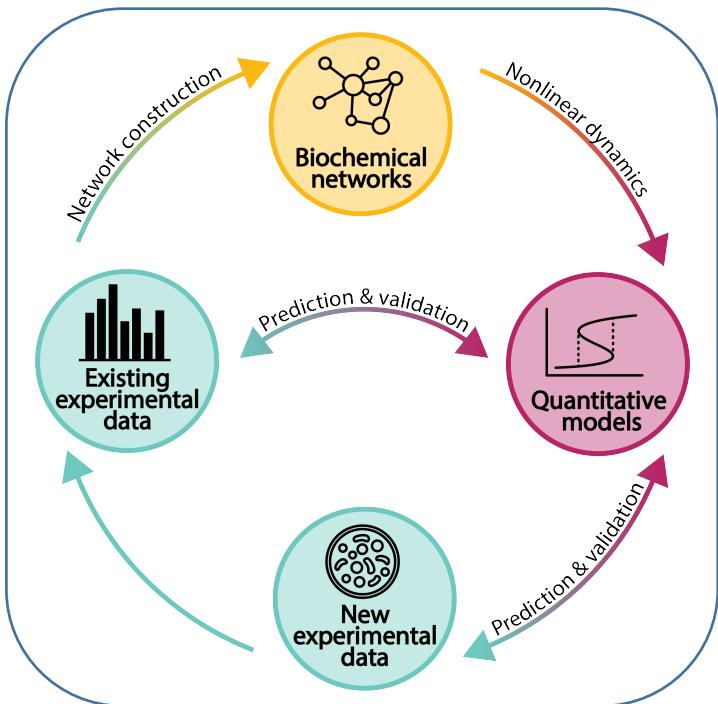


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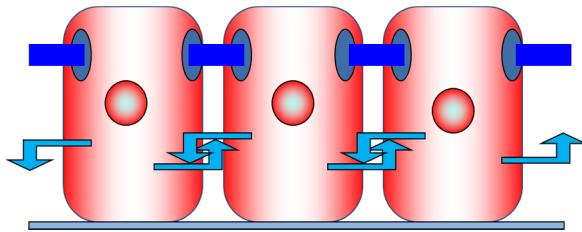
MHRD



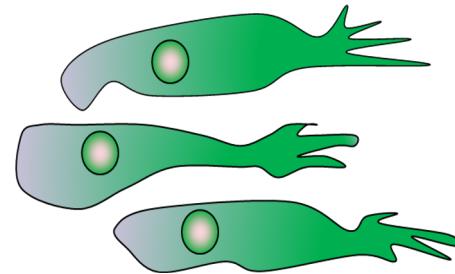
Department of Science & Technology
Government of India



EMT/MET: The engine of metastasis



Adhere to neighbors
Do NOT migrate or invade
Epithelial (E)

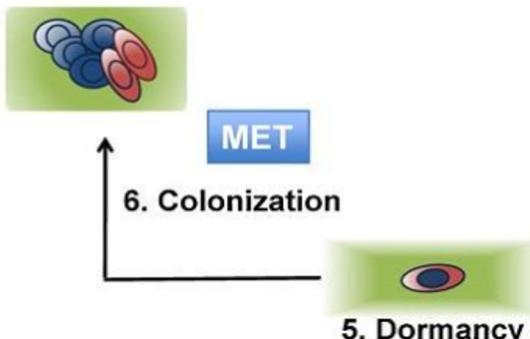


Do NOT adhere to neighbors
Migrate and invade
Mesenchymal (M)

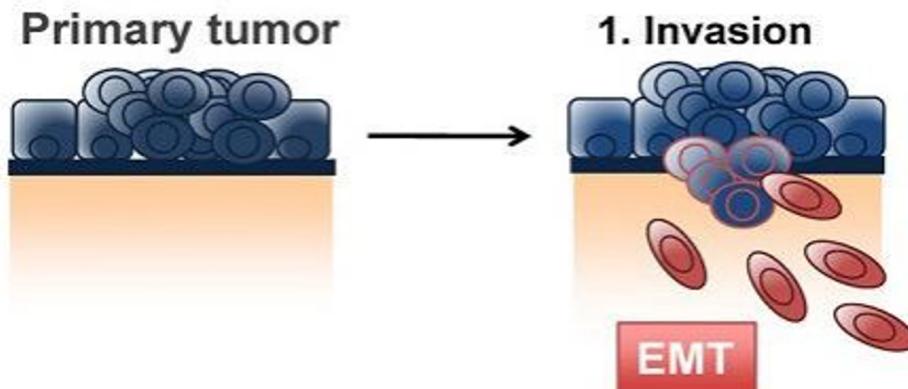


Mesenchymal-to-Epithelial
Transition (MET)

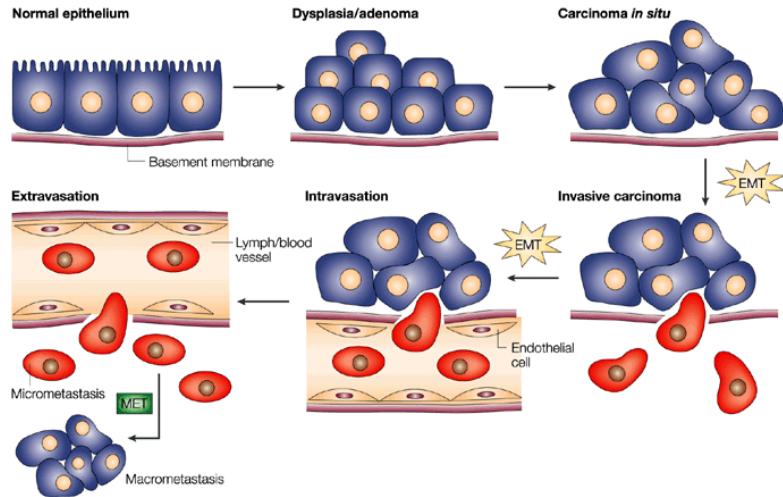
Secondary tumor



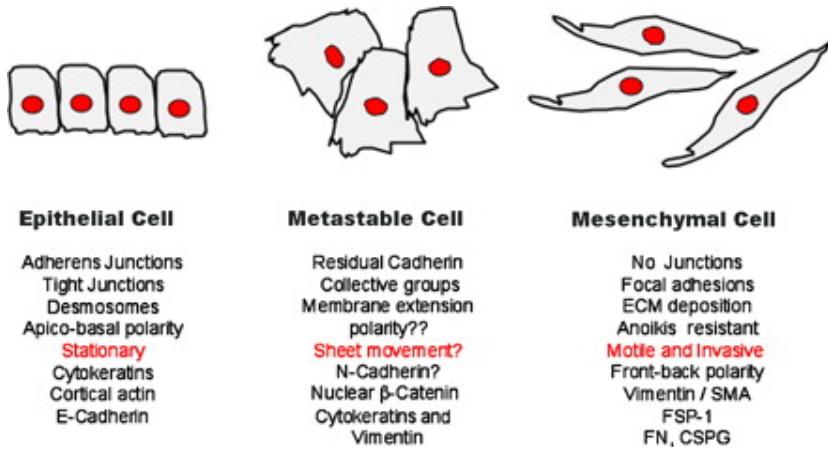
Epithelial-to-Mesenchymal
Transition (EMT)



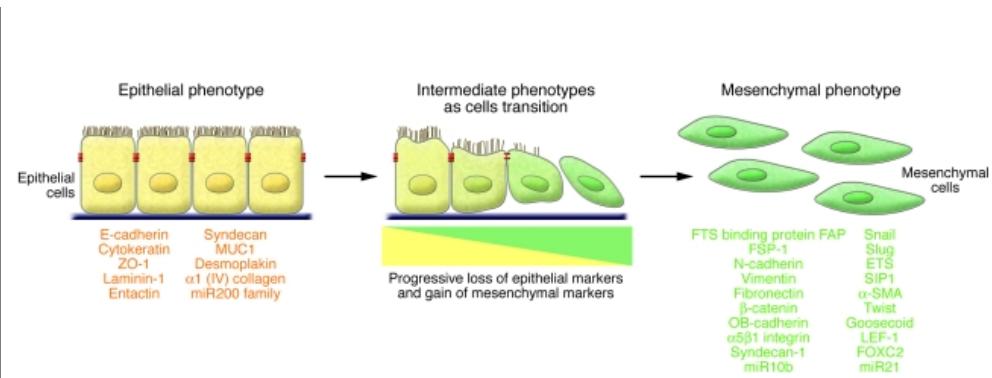
Role of EMT in cancer metastasis (2002 – 2012)



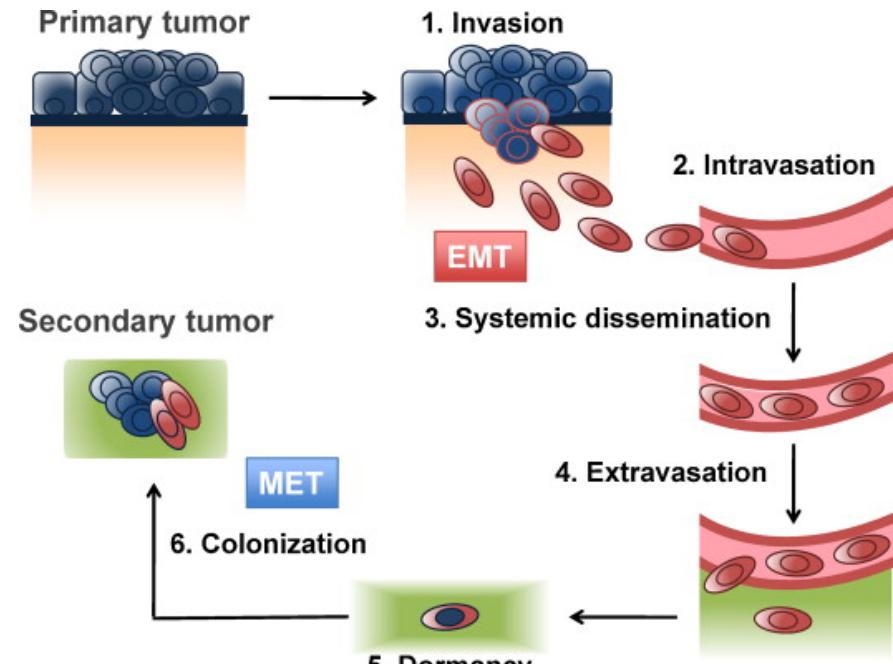
Thiery JP, Nat Rev Cancer 2002



Lee et al. J Cell Biol 2006

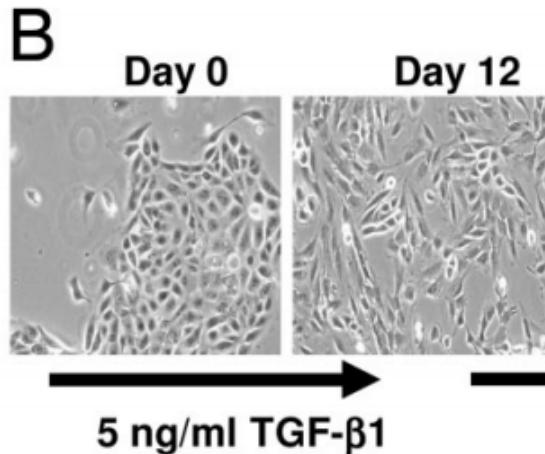


Kalluri & Weinberg, J Clin Invest 2009



Scheel & Weinberg, Semin Cancer Biol 2012

Is EMT/MET a binary process?

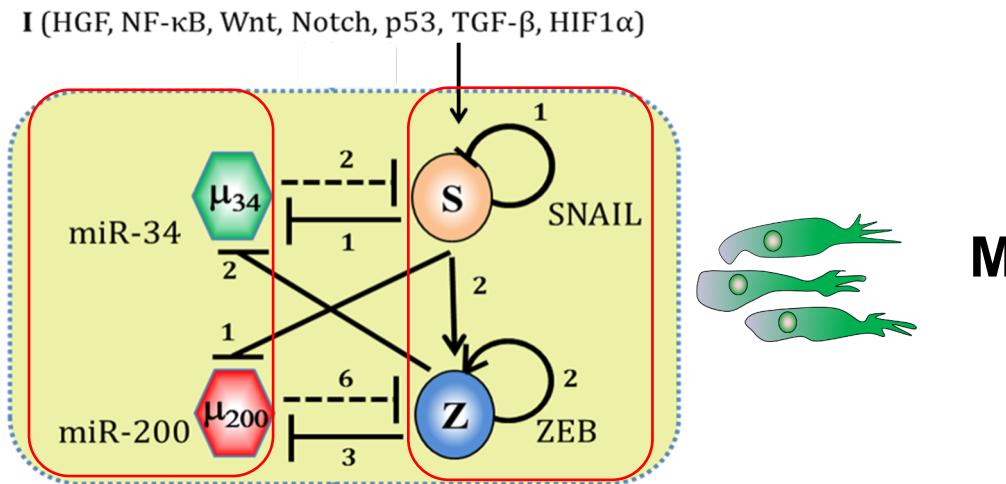
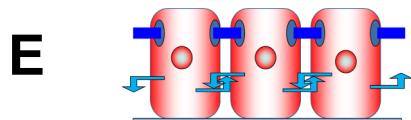


Mani et al.
PNAS 2007



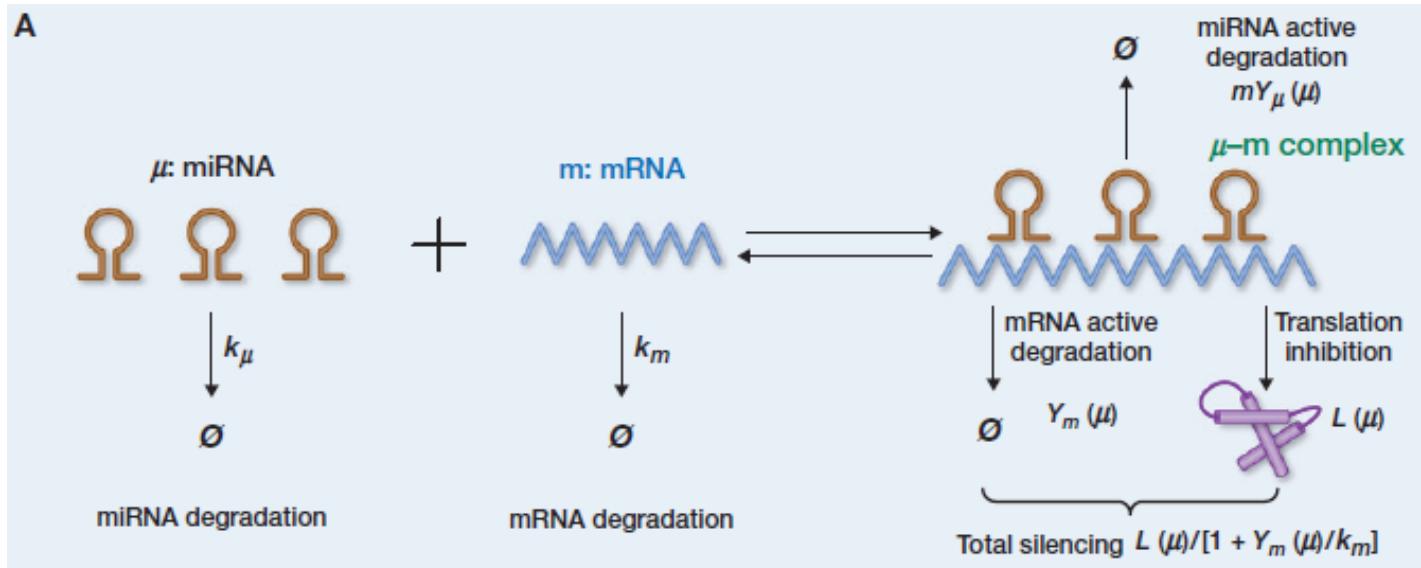
Network that controls EMT/MET

→ Transcriptional activation
→| Transcriptional repression
---+ miR-mediated repression



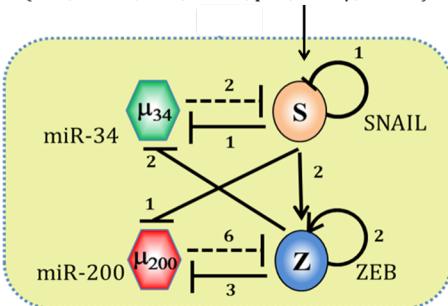
- Each arrow/bar indicates a quantitative input-output relationship.
- Such models have been extensively built for simpler microorganisms.
- Can we decode the emergent properties of these nonlinear interactions?

Mathematical model formulation



Lu*, Jolly* et al. PNAS 2013

I (HGF, NF-κB, Wnt, Notch, p53, TGF-β, HIF1α)



Production

Degradation

miR regulation

TF regulation

$$\frac{d\mu_{200}}{dt} = g_{\mu_{200}} H^S(Z, \lambda_{Z,\mu_{200}}) H^S(S, \lambda_{S,\mu_{200}}) - m_Z Y_\mu(\mu_{200}) - k_{\mu_{200}} \mu_{200}$$

miR-200

$$\frac{dm_Z}{dt} = g_{m_Z} H^S(Z, \lambda_{Z,m_Z}) H^S(S, \lambda_{S,m_Z}) - m_Z Y_m(\mu_{200}) - k_{m_Z} m_Z$$

ZEB mRNA

$$\frac{dZ}{dt} = g_Z m_Z L(\mu_{200}) - k_Z Z$$

ZEB

$$\frac{d\mu_{34}}{dt} = g_{\mu_{34}} H^S(Z, \lambda_{Z,\mu_{34}}) H^S(S, \lambda_{S,\mu_{34}}) - m_S Y_\mu(\mu_{34}) - k_{\mu_{34}} \mu_{34}$$

miR-34

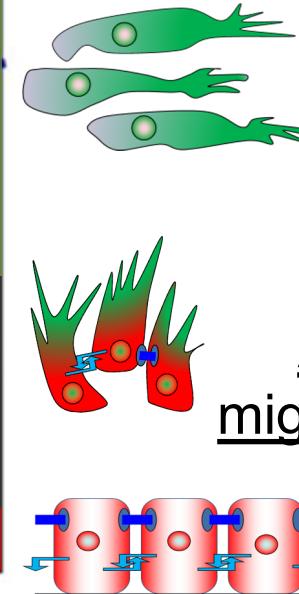
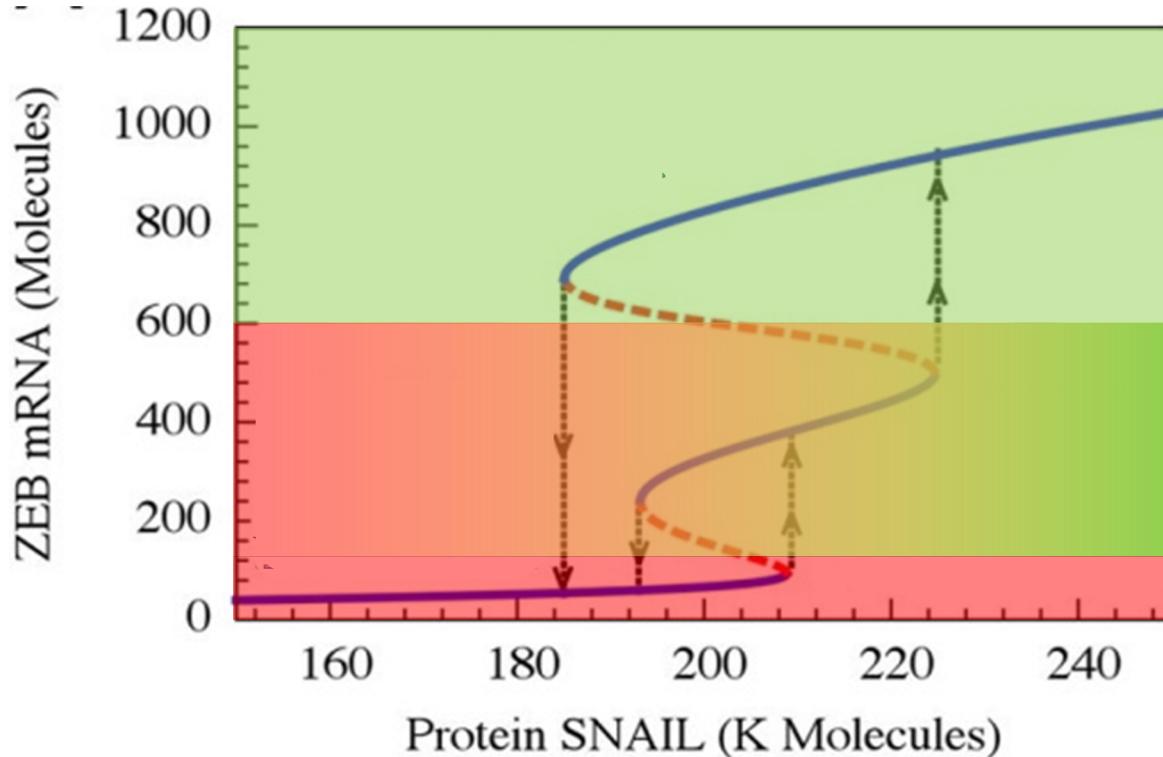
$$\frac{dm_S}{dt} = g_{m_S} H^S(S, \lambda_{S,m_S}) H^S(I, \lambda_{I,m_S}) - m_S Y_m(\mu_{34}) - k_{m_S} m_S$$

SNAIL mRNA

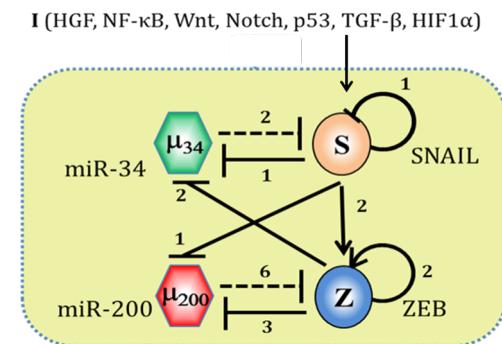
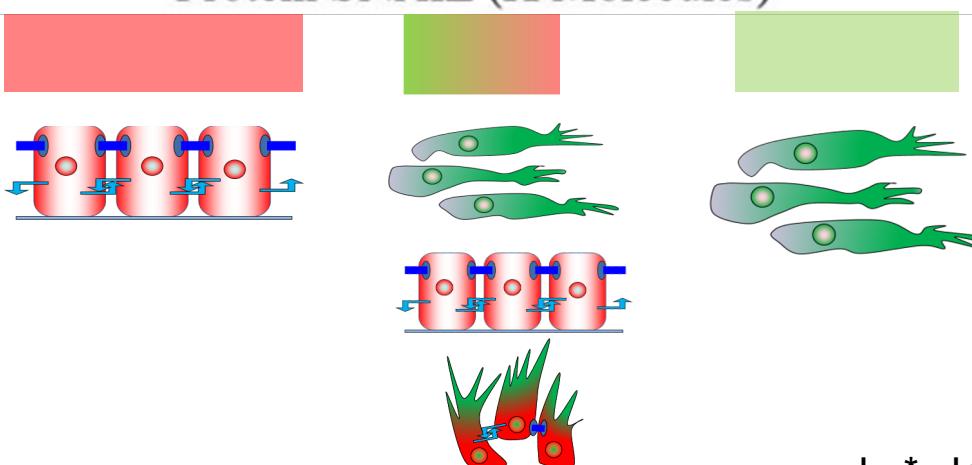
$$\frac{dS}{dt} = g_S m_S L(\mu_{34}) - k_S S$$

SNAIL₁₂

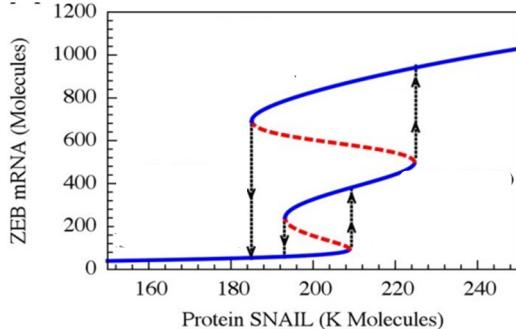
Model prediction: EMT is NOT binary



Hybrid E/M
Adhere AND
migrate collectively



Mathematical modeling for EMT dynamics

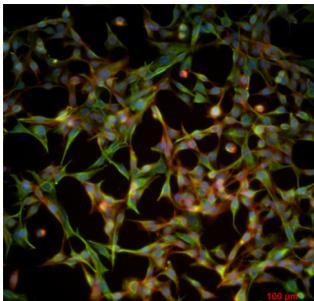


Predictions from mathematical model:

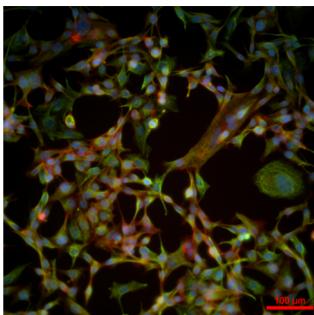
1. Cells can stably exist in hybrid E/M state
2. Isogenic cells can exist in different EMT states
3. Cells can ‘spontaneously’ switch their states

Lu*, Jolly* et al. PNAS 2013

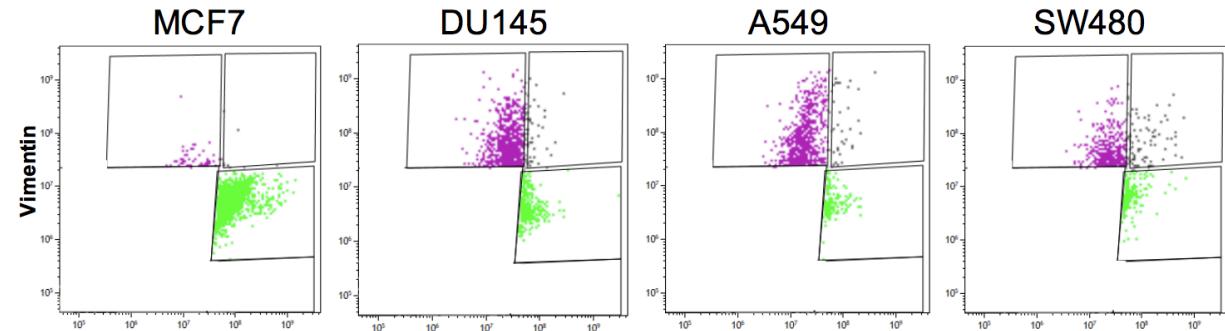
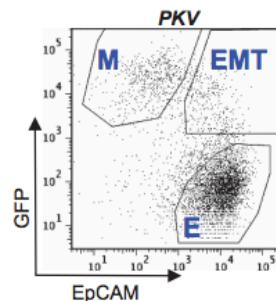
Experimental validation:



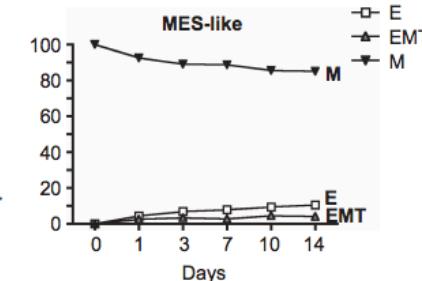
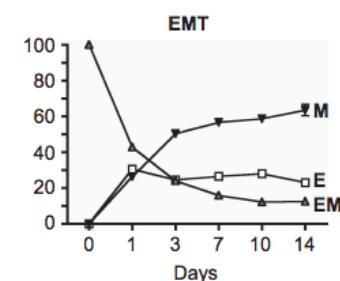
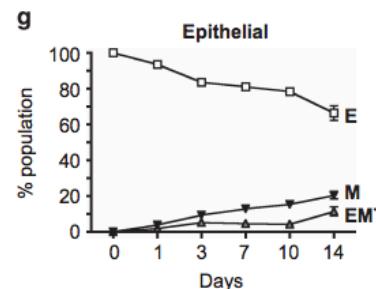
H1975, T=0



H1975, T=2 months



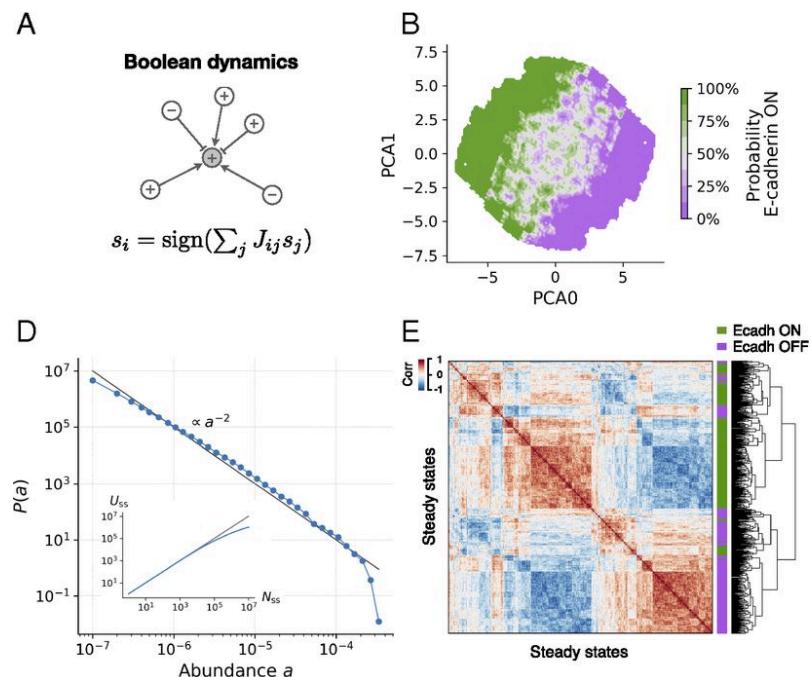
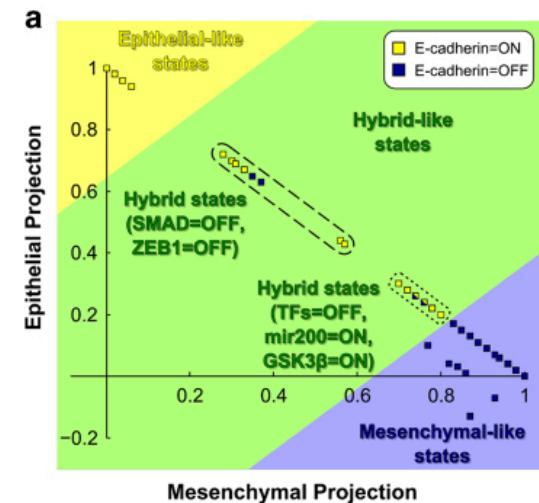
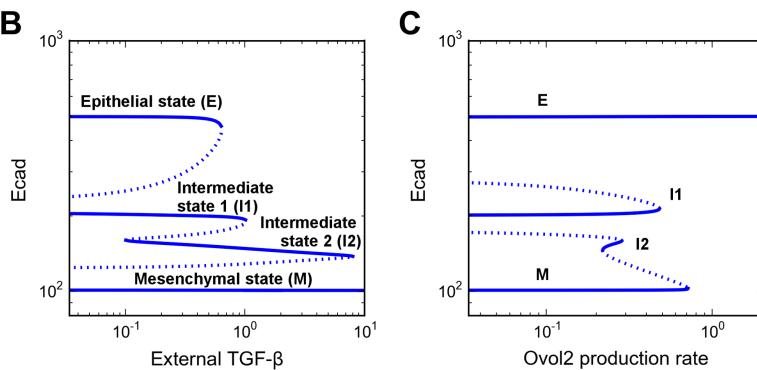
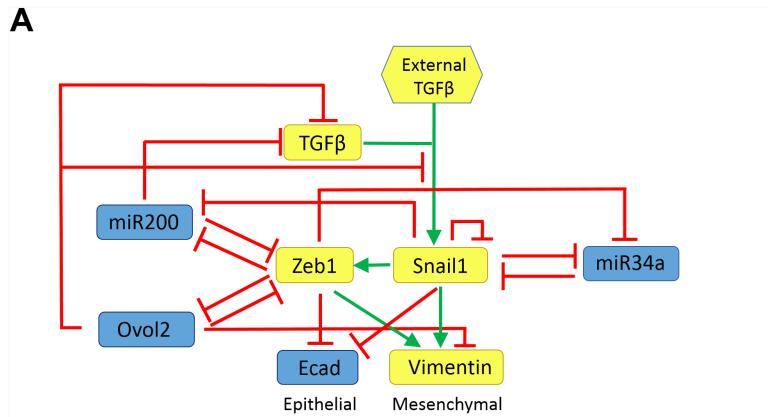
George*, Jolly* et al. Cancer Res 2017



Jolly et al. Oncotarget 2016

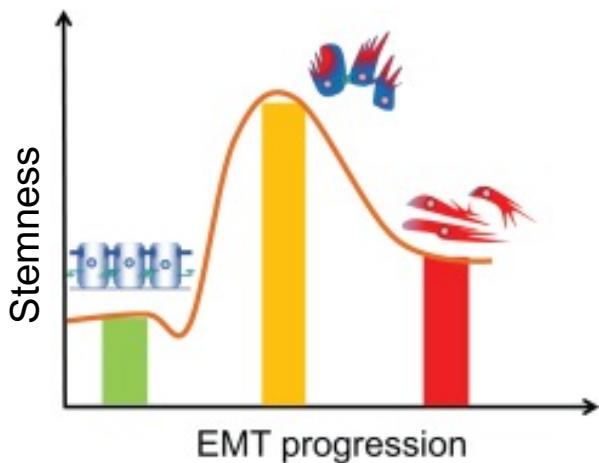
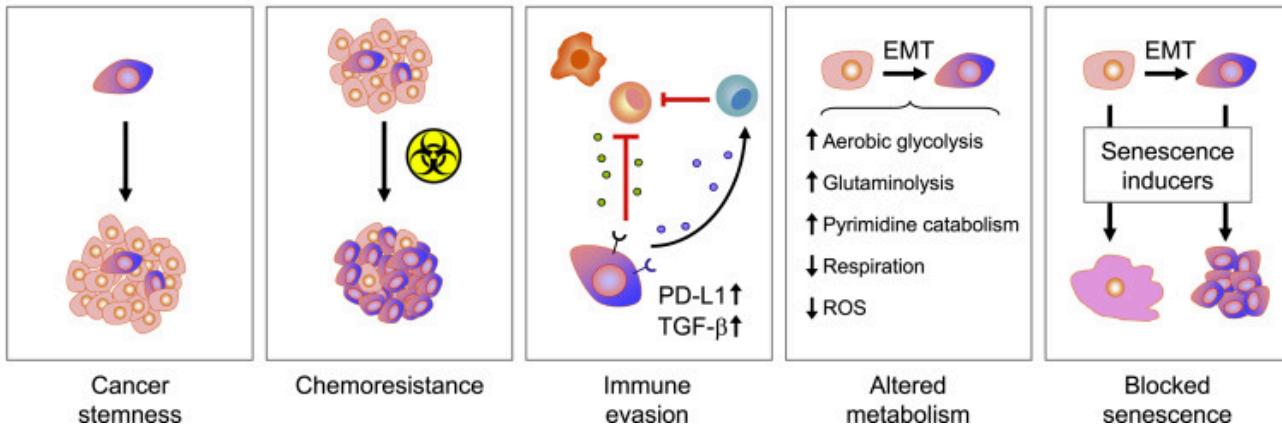
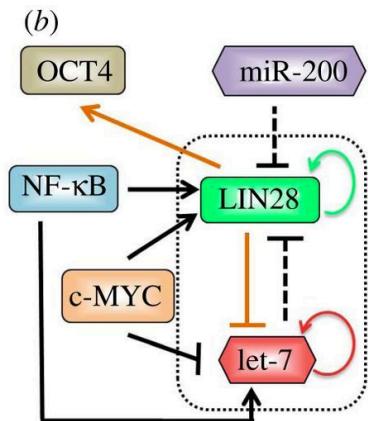
Ruscetti et al. Oncogene 2016

Hybrid E/M phenotype(s) seen in other math models too



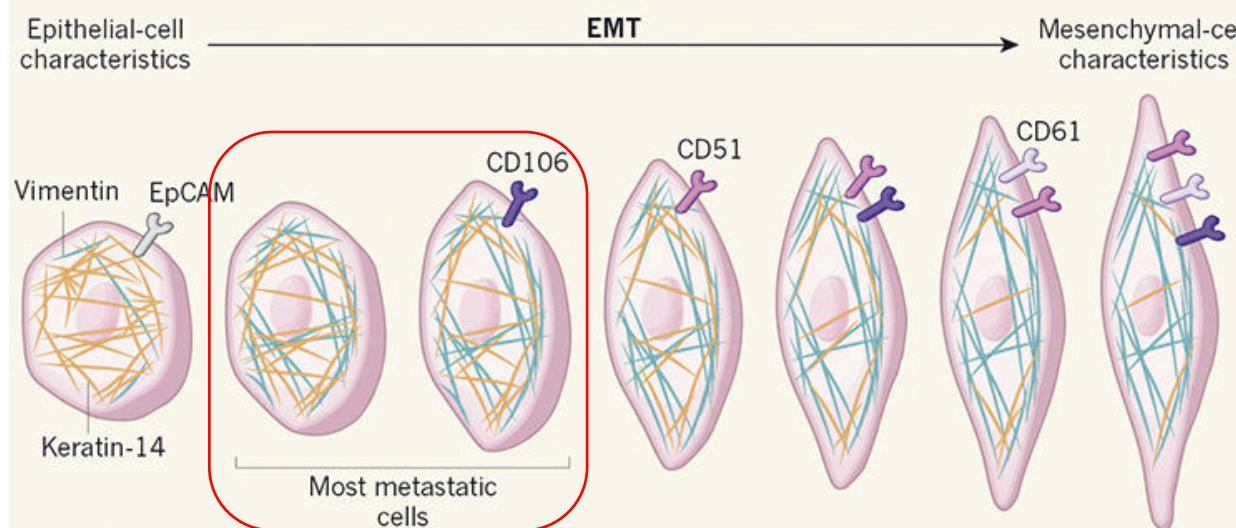
- Xing *et al.* Biophys J 2013
 Steinway *et al.* NPJ Sys Biol Appl 2015
 Hong *et al.* PLoS Comp Biol 2015
 Jolly *et al.* Oncotarget 2016
 Huang *et al.* PLoS Comp Biol 2017
 Font-Clos *et al.* PNAS 2018
 Silveira *et al.* FEBS J 2019
 Hari *et al.* NPJ Sys Biol Appl 2020

Hybrid E/M phenotype(s): ‘fittest’ for metastasis?



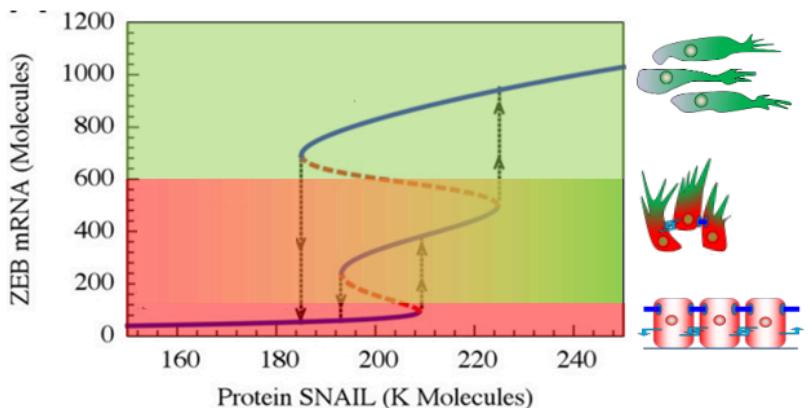
Acquisition of a hybrid E/M state is essential for tumorigenicity of basal breast cancer cells

Cornelia Kröger^a, Alexander Afeyan^{a,b}, Jasmin Mraz^{a,c}, Elinor Ng Eaton^a, Ferenc Reinhardt^a, Yevgenia L. Khodor^d, Prathapan Thiru^a, Brian Bierie^a, Xin Ye^{a,e}, Christopher B. Burge^d, and Robert A. Weinberg^{a,f,g,1}

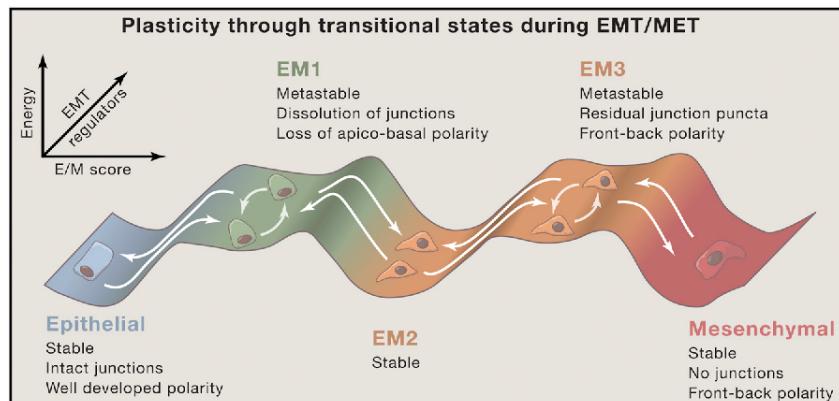


- Jolly et al. J R Soc Interface 2014
Grosse-Wilde et al. PLoS One 2015
Bierie et al. PNAS 2017
Pastushenko et al. Nature 2018
Kroger et al. PNAS 2019
Lu & Kang, Dev Cell 2019
Pastushenko et al. Nature 2021

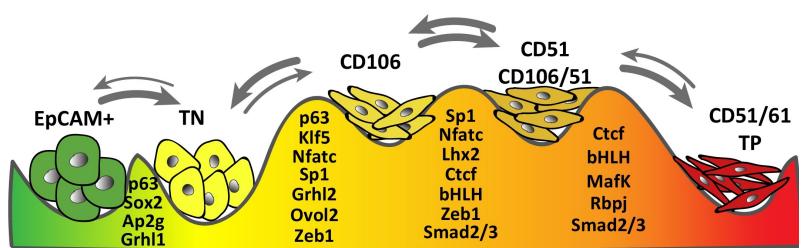
From EMT (2002-2012) to EMP (Epi-Mes Plasticity; 2013-now)



Lu[#], Jolly[#] et al. PNAS 2013

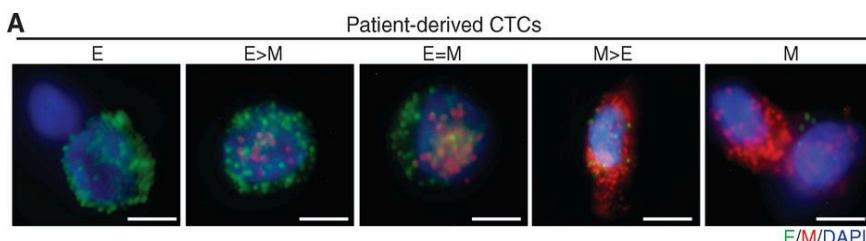


Nieto et al. Cell 2016



	Epithelial tumor cells	Early hybrid EMT state	Hybrid EMT state	Late hybrid EMT state	Mesenchymal tumor cells
Proliferation	++++	+++	++	++	+
Invasion	+	++	+++	++++	+++++
Plasticity	+	++	+++	++++	++
Stemness	+	+++	+++	+++	+++
Metastasis	+	++++	++++	++	+

Pastushenko & Blanpain, Trends Cell Biol 2019
Pastushenko et al. Nature 2018

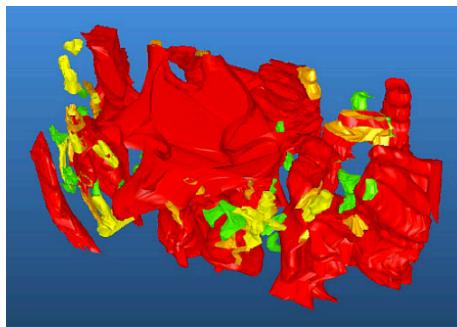
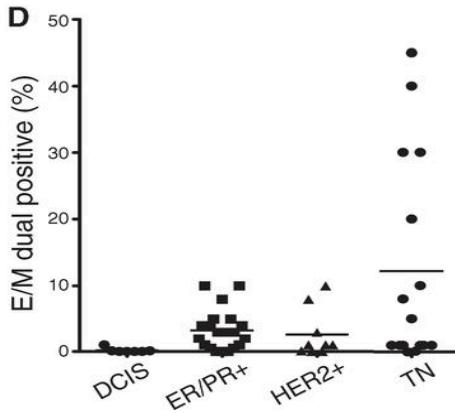


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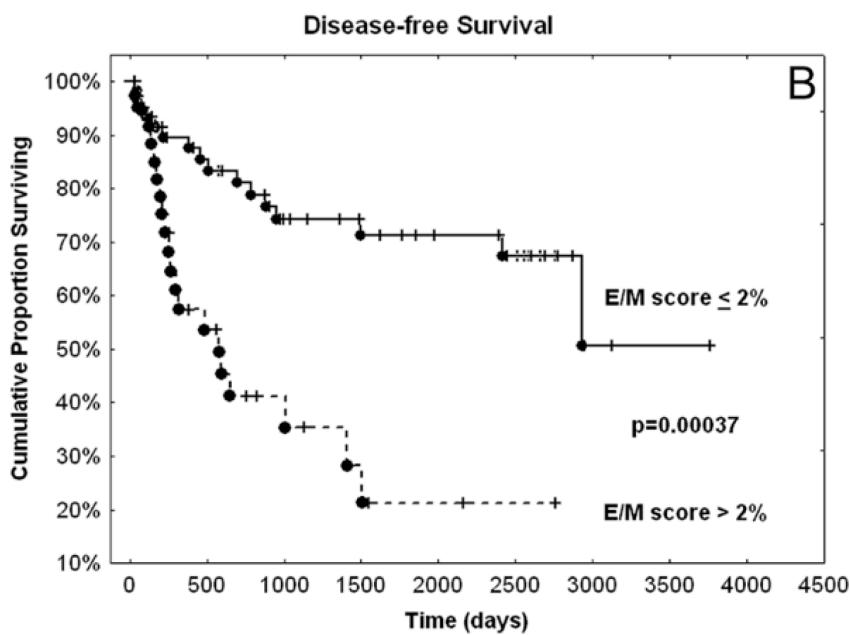
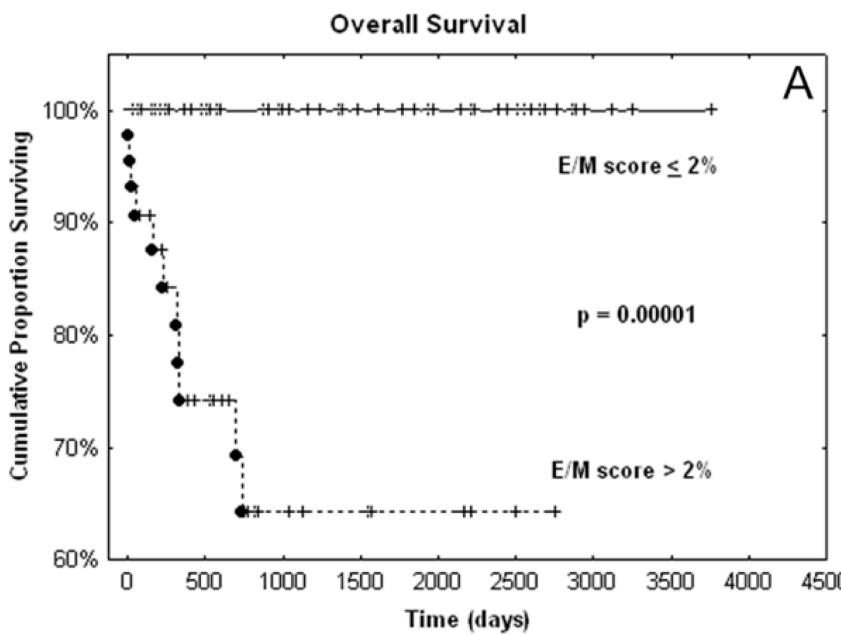
Yu et al. Science 2013
Kroger et al. PNAS 2019

Clinical relevance of hybrid E/M phenotype(s)



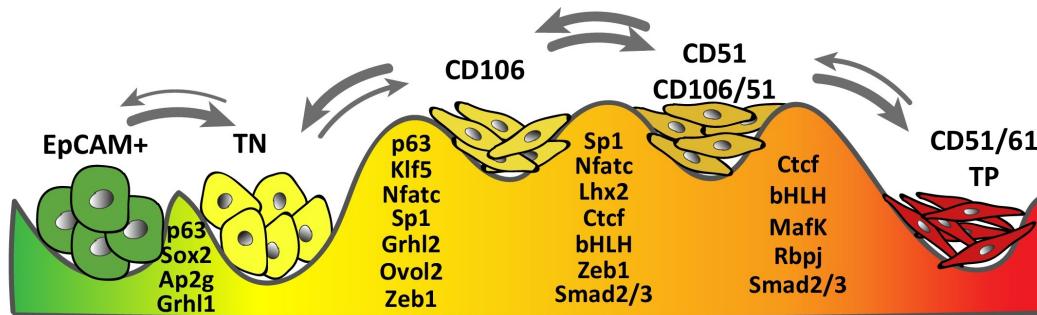
Single-cell migration is very rare, if any, in cancer

Co-expression of ZEB1 and membranous E-cad - a 'partial EMT' status of 'tumor buds'



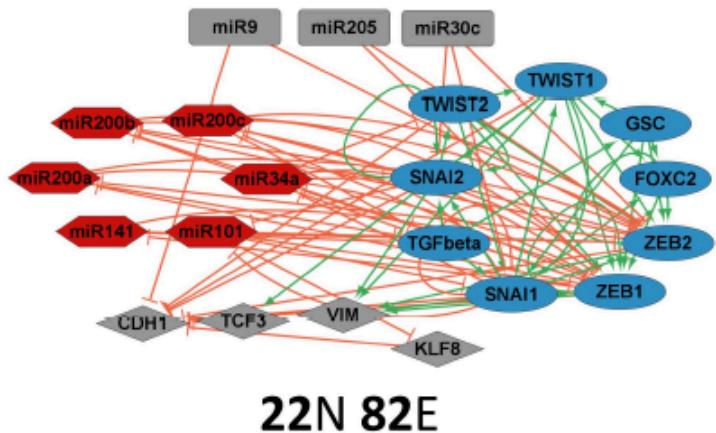
Ongoing questions...

Why are hybrid E/M cells more plastic than E, M?

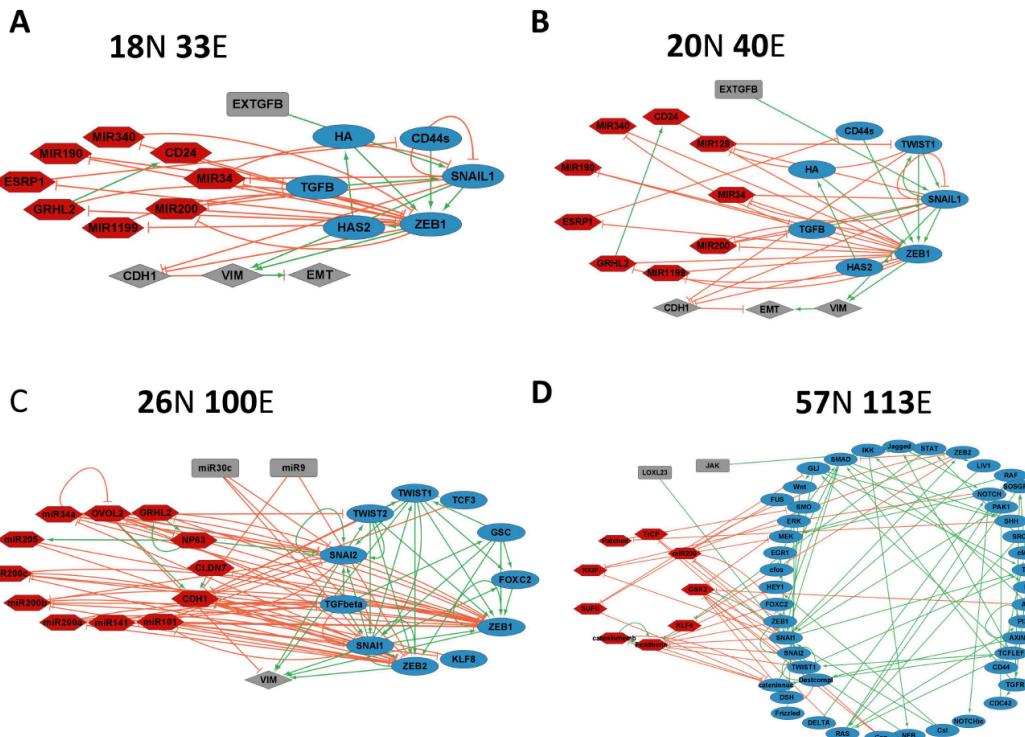


	Epithelial tumor cells	Early hybrid EMT state	Hybrid EMT state	Late hybrid EMT state	Mesenchymal tumor cells
Proliferation	+++++	+++	++	++	+
Invasion	+	++	++	+++	++++
Plasticity	+	++	++	+++	++
Stemness	+	++	++	++	++
Metastasis	+	+++	+++	++	+

EMP networks largely contain two “teams”

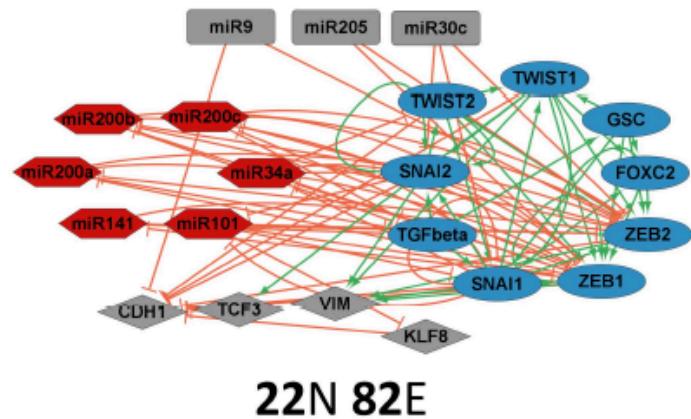


Huang *et al.* PLoS Comp Biol 2017
Jia *et al.* Phys Biol 2019
Font-Clos *et al.* PNAS 2018
Silveira *et al.* FEBS J 2019; JRSI 2020



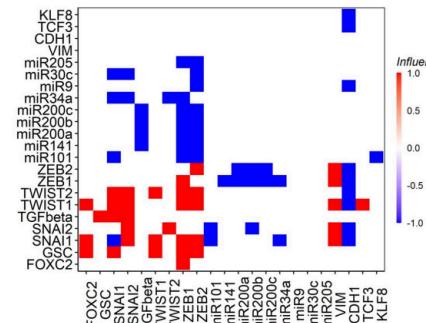
- Nodes: **Epithelial**, **Mesenchymal**, Input/output (Signal/Effectors)
- Edges: **Activation**, **Inhibition**
- Mostly activation within a “team”, but inhibition across the two “teams”

Presence of teams is specific to EMP networks

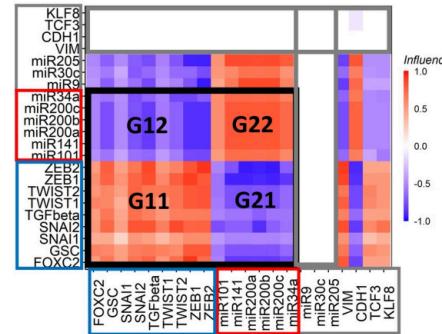


22N 82E

22N 82E
Adjacency Matrix



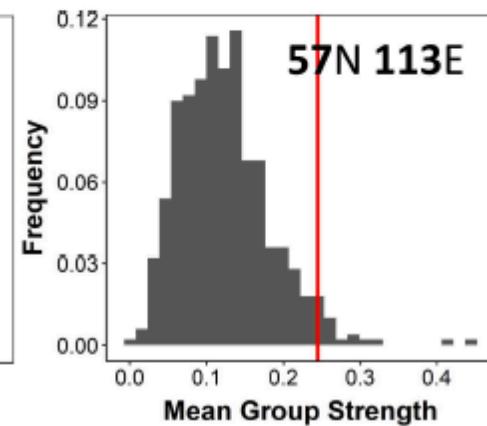
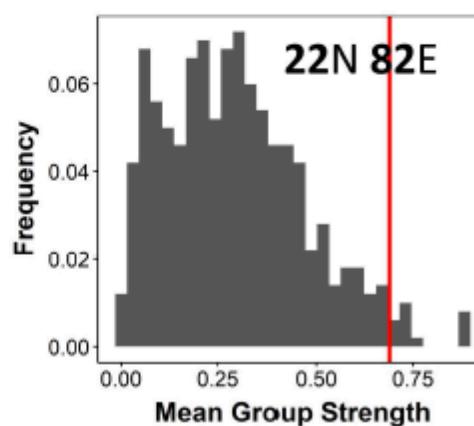
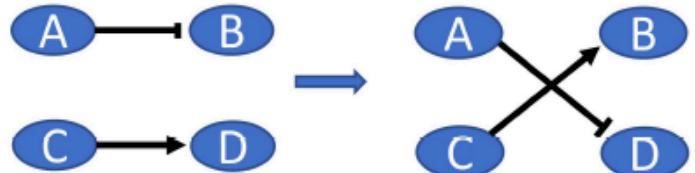
22N 82E
Influence Matrix



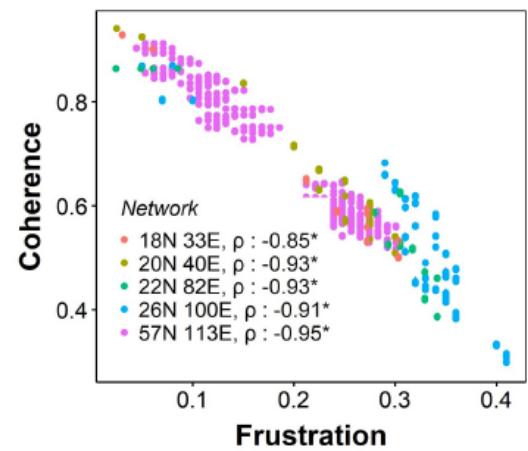
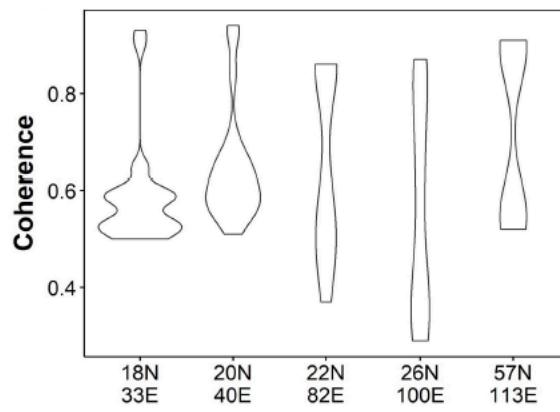
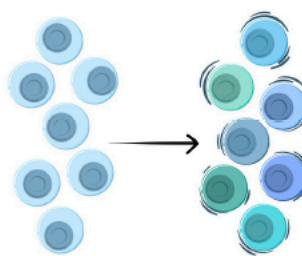
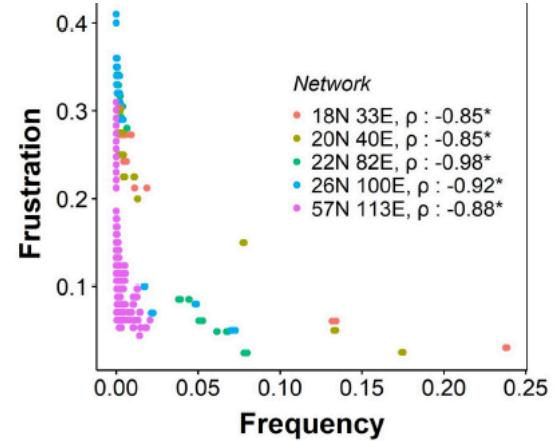
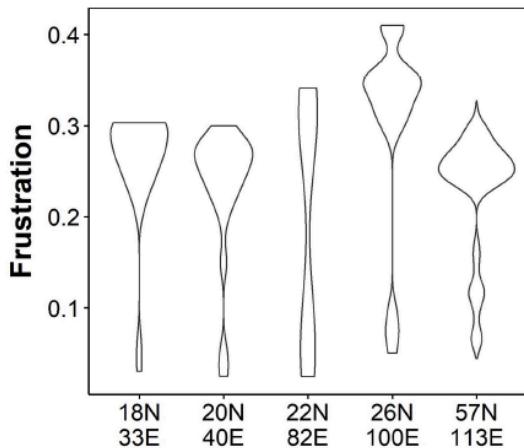
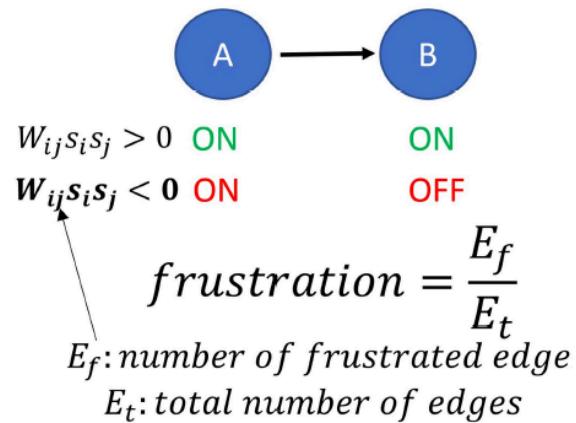
$$Infl = \frac{\sum_{l=1}^{lmax} \frac{Adj^l}{Adj_{max}}}{lmax}$$

$$G_{IJ} = \frac{\sum_{i \in I, j \in J} Inf_{ij}}{n_{II}}$$

$$G_S = \sum_{i,j \in \{1,2\}} |G_{ij}|$$

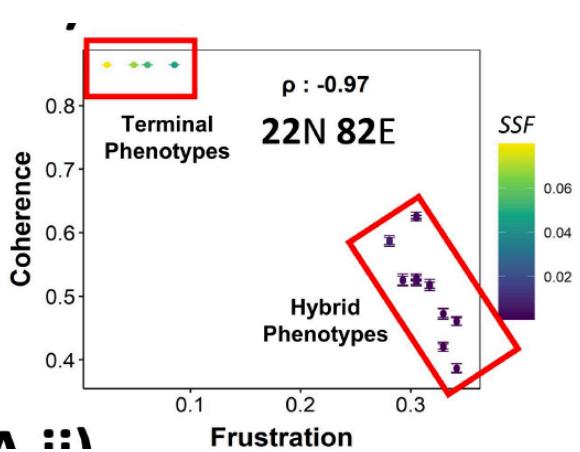


EMP networks give rise to two types of states

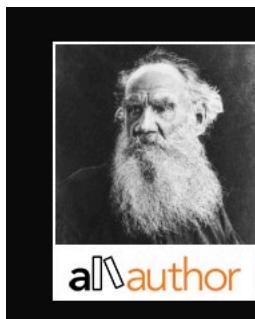
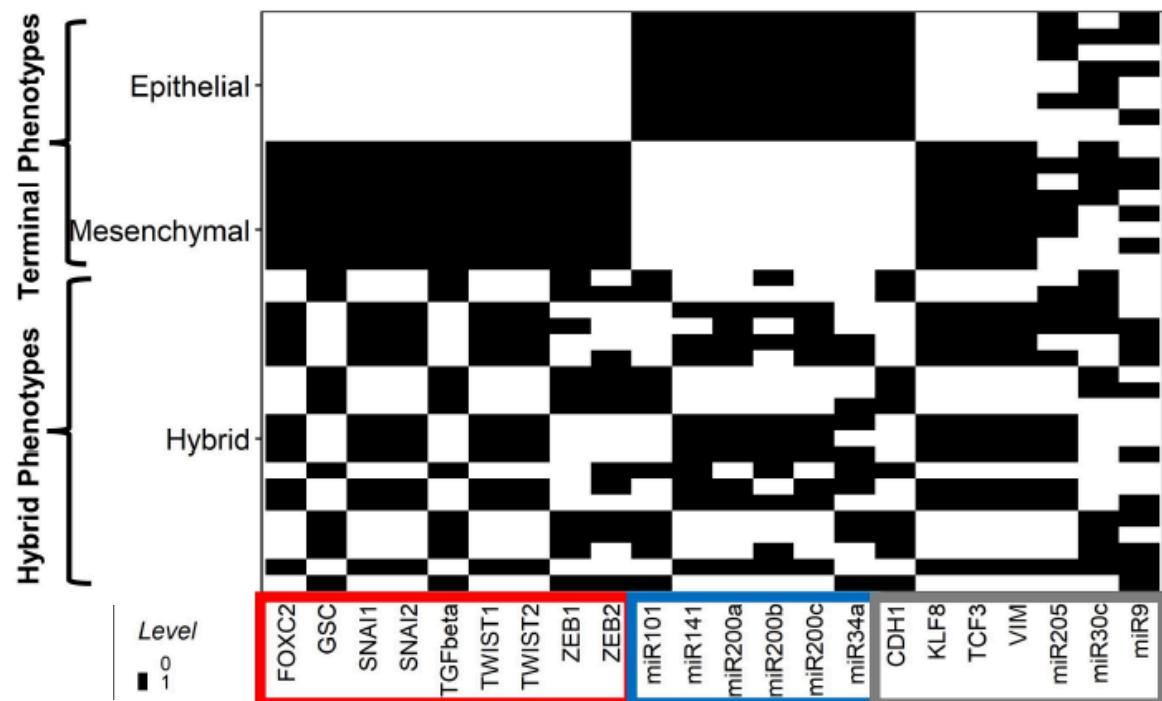


Which phenotypes (E, M, hybrid) are more frustrated or coherent?

Terminal states (E, M) more stable than hybrid E/M



Terminal state more
coherent, frequent; less
frustrated than hybrid



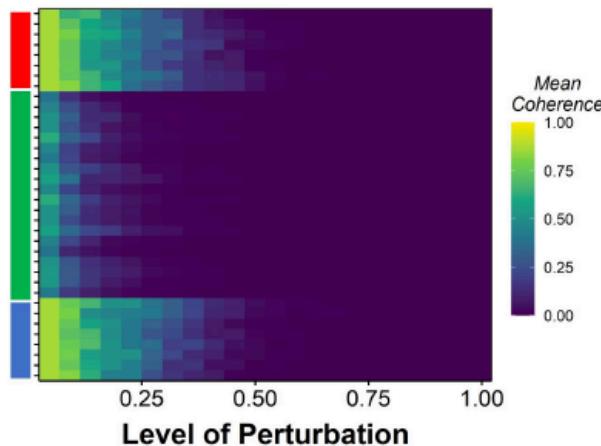
All happy families are alike; each unhappy family is unhappy in its own way.

-Leo Tolstoy

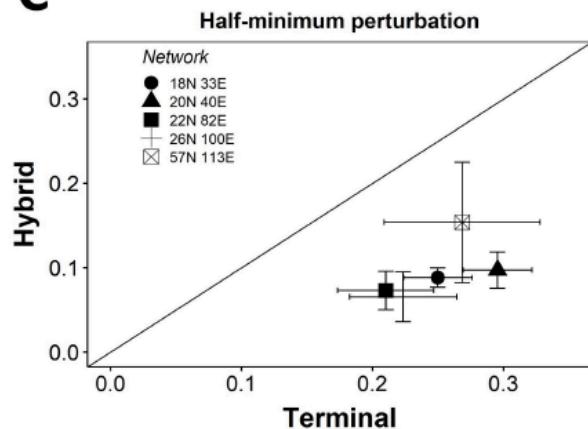
“Teams” stabilize terminal states (E, M) specifically

B

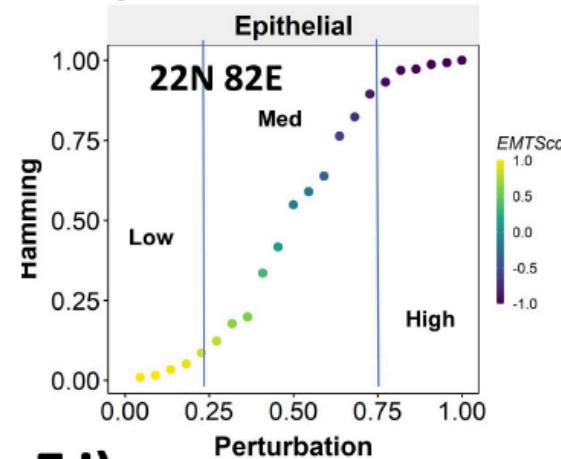
22N 82E



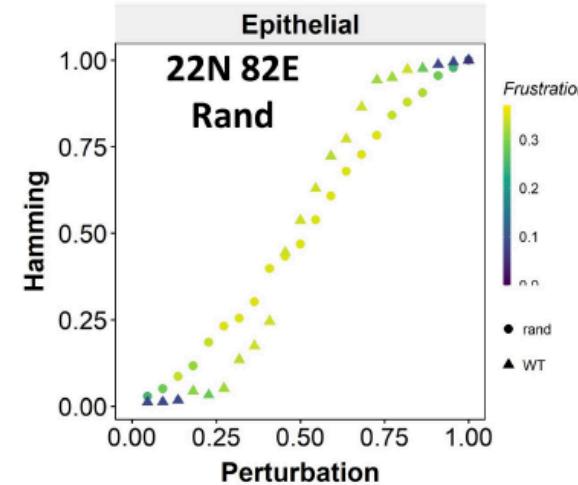
C



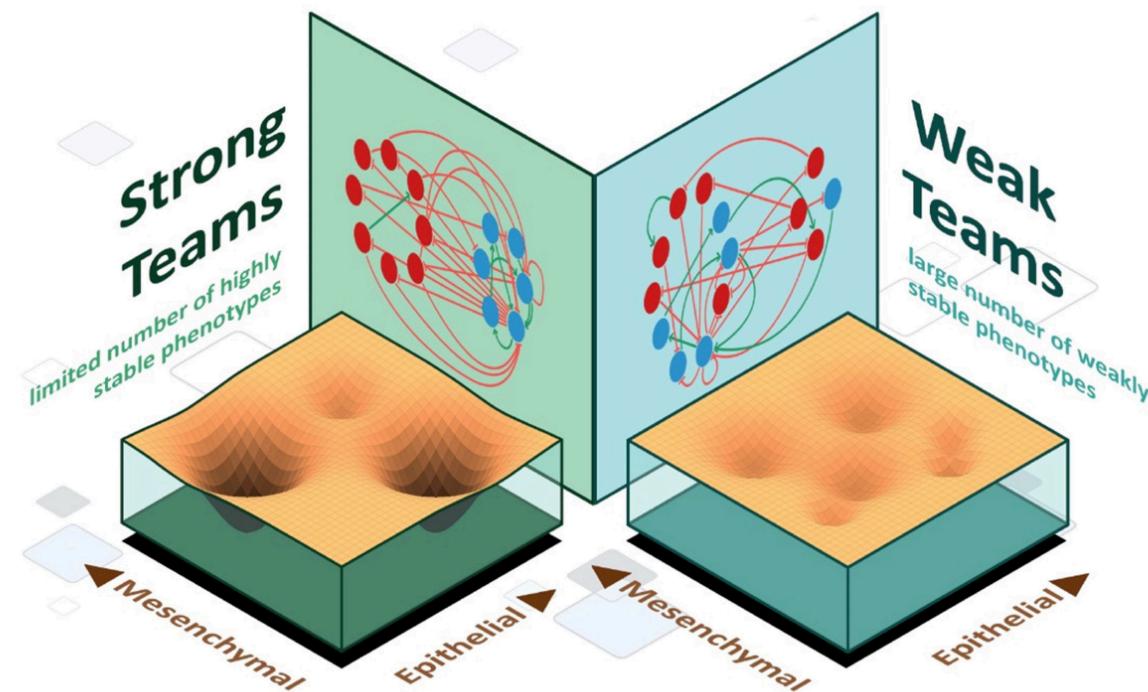
D i)



E i)

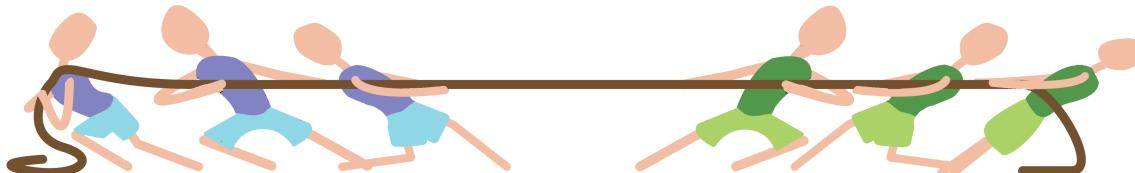


Summary (Teams in EMP networks)



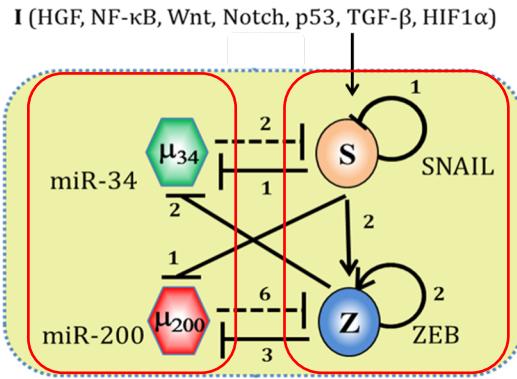
“Teams” shape the landscape enabling higher plasticity and heterogeneity of hybrid E/M phenotypes

Hari *et al.* bioRxiv 2021: 472090

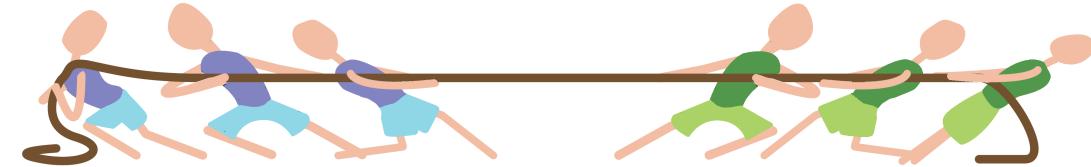


Kishore Hari

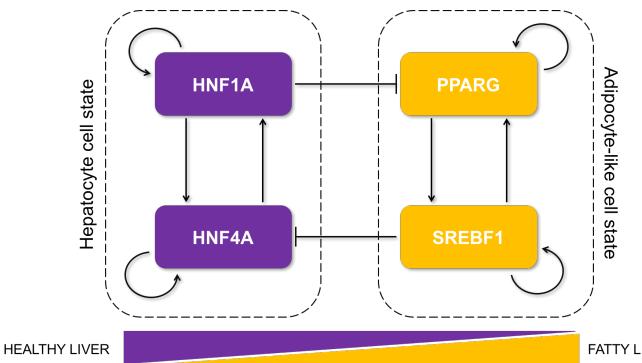
'Teams' seen in other cell-state switching networks?



Lu*, Jolly* et al. PNAS 2013



Artwork Credit: Atchuta S Duddu



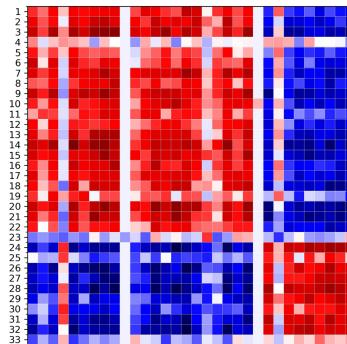
Sahoo*, Singh* et al. J Clin Med 2020



Divyoj Singh
(BS/MS,
IISc, Phy)

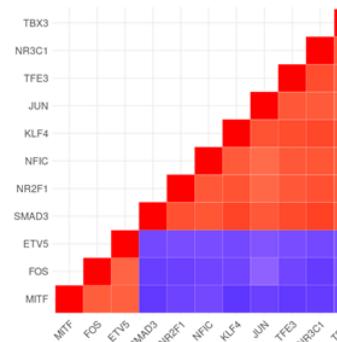


Sarthak Sahoo
(BS/MS,
IISc, Bio)



**Small Cell Lung
Cancer (SCLC)**

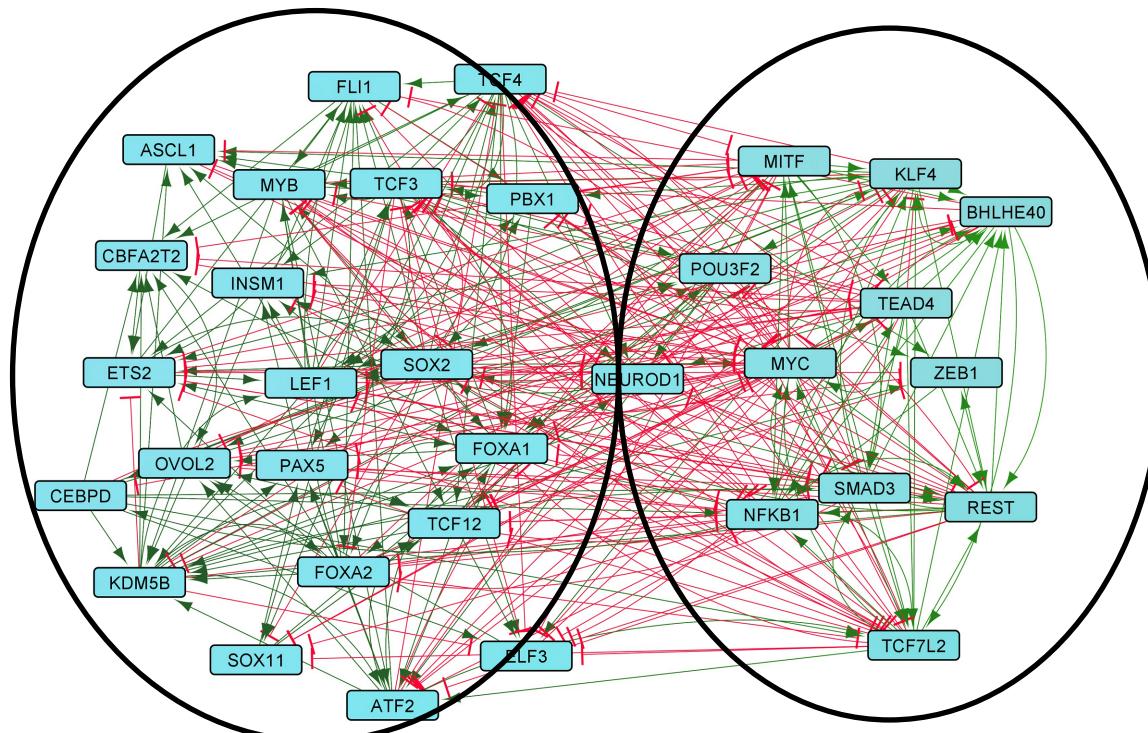
Chauhan*, Ram*
et al. eLife 2021



**Proliferative-invasive
switch in melanoma**

Pillai & Jolly, iScience 2021

Why do ‘teams’ exist?

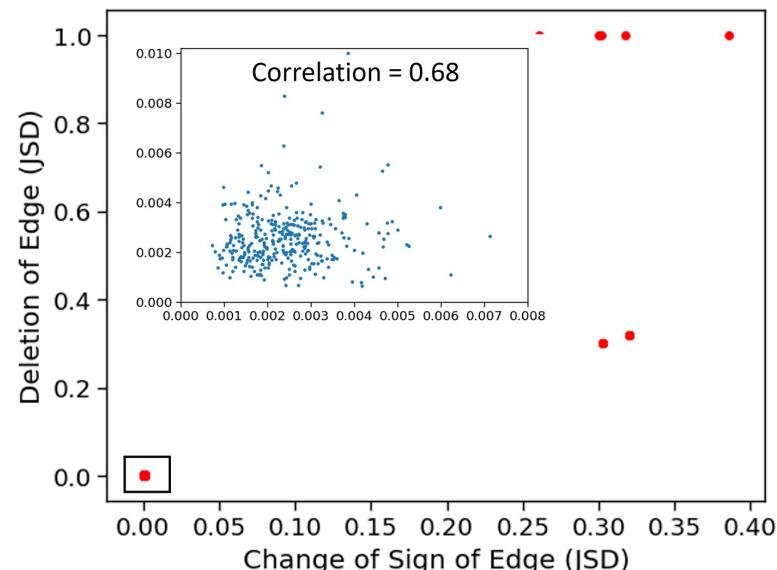


33 nodes, 357 edges - **SCLC**

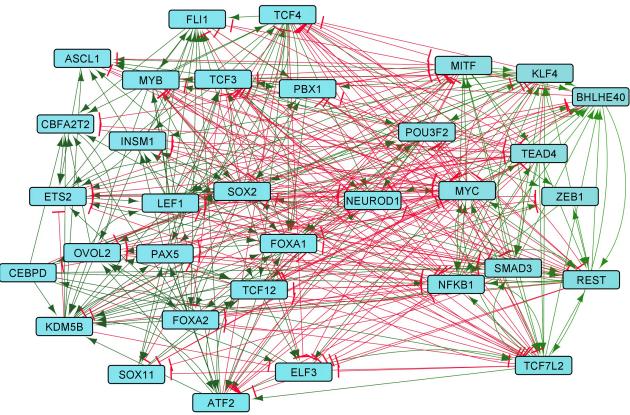
Offer robustness (Deletion of 345 out of 357 edges – one at a time – have minimal effect on SCLC cell states)

This network gave rise to 4 states, each with ~25% frequency.

What if we delete an edge (in silico CRISPR)?



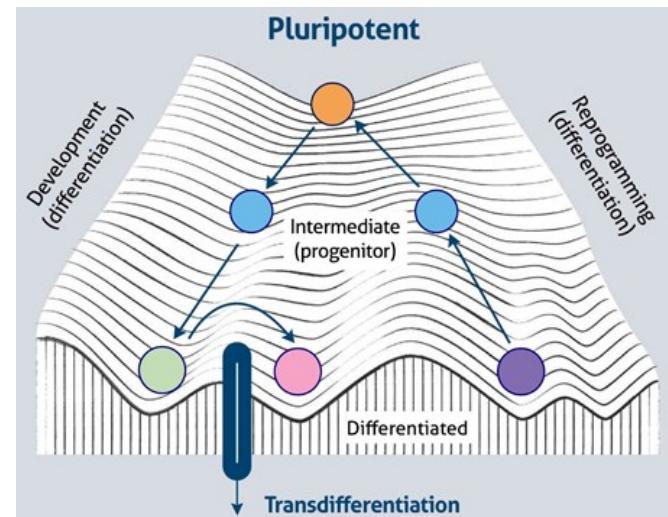
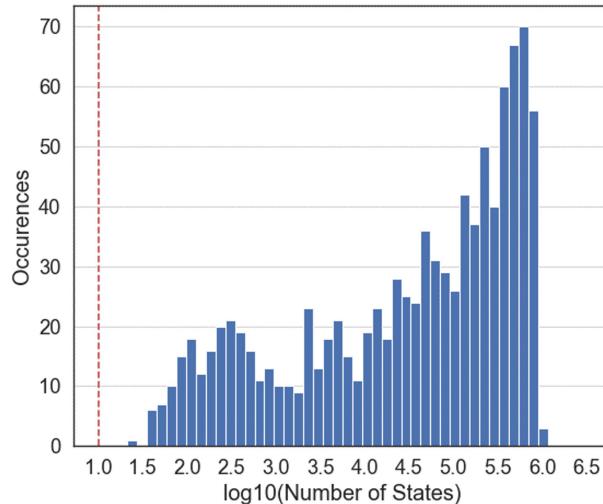
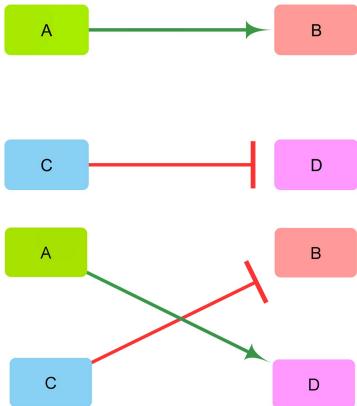
Why do ‘teams’ exist?



33 nodes, 357 edges - SCLC



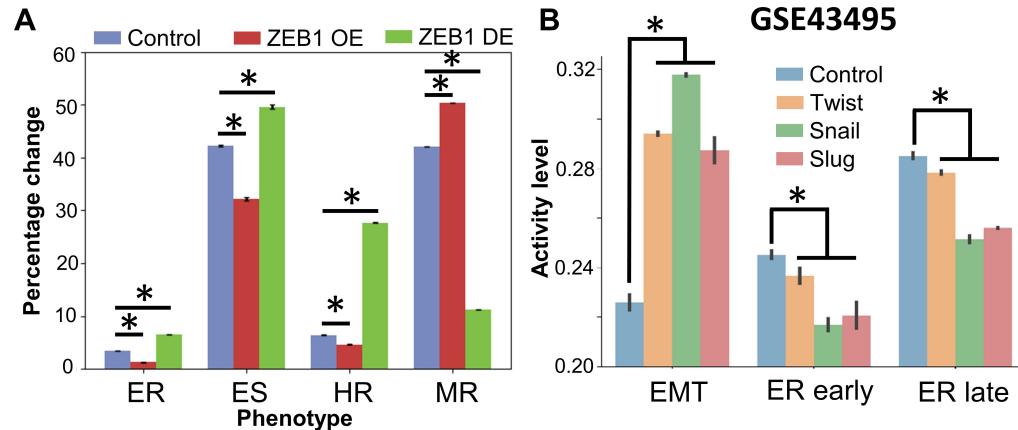
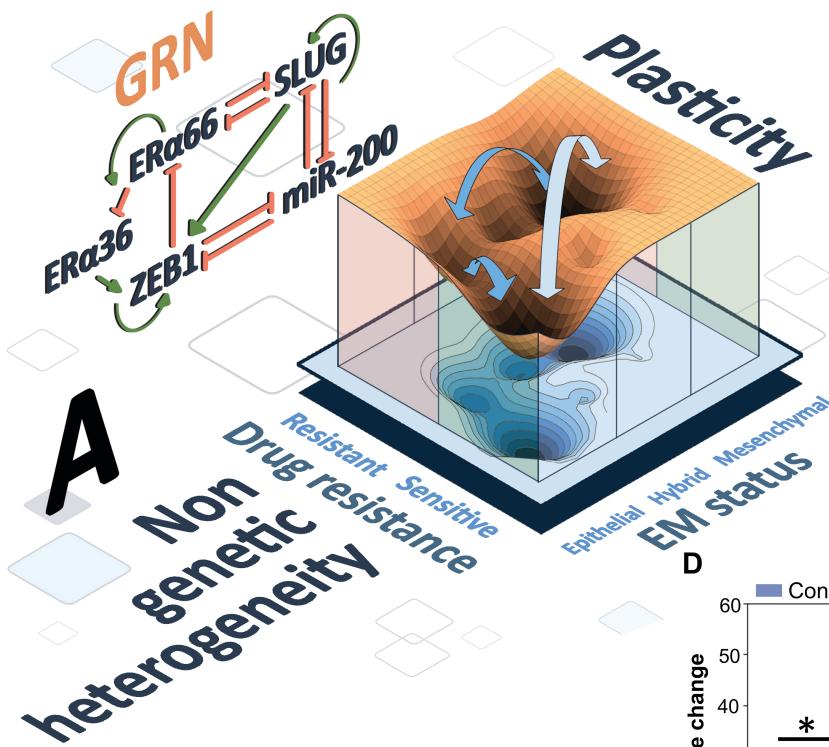
What if we shuffle edges in the entire network (thus breaking teams)?



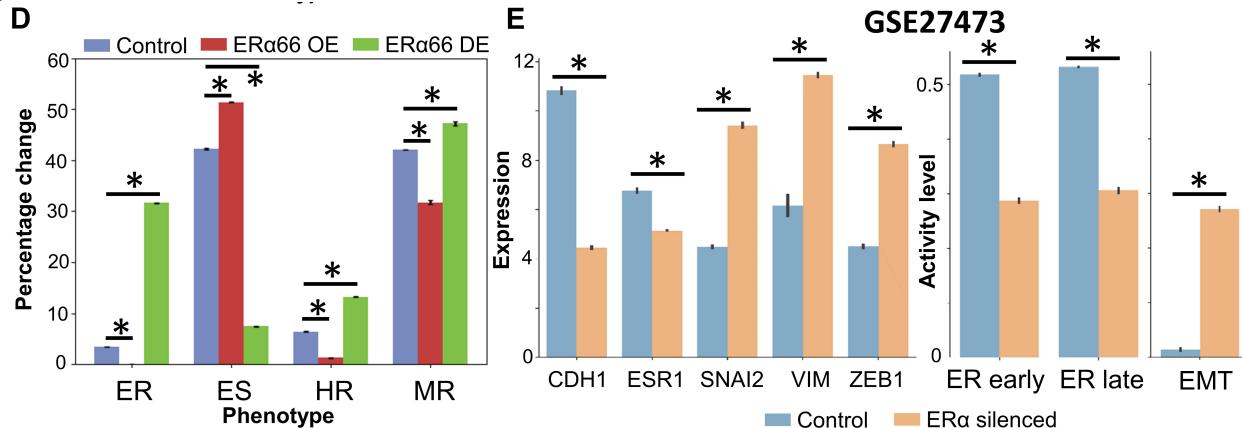
Allow limited number of cell-states
("Controlled enthusiasm")

Why do ‘teams’ exist?

Couple the axes of plasticity: EMP and drug resistance in ER+ breast cancer



EMT can drive resistance to tamoxifen



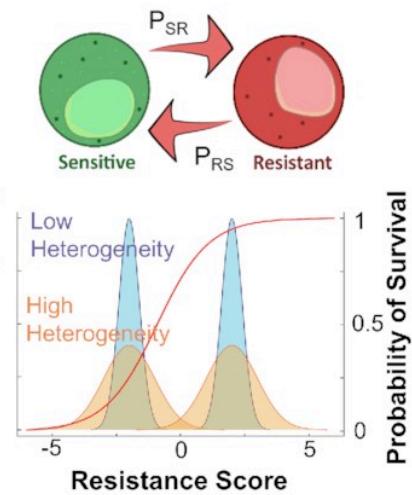
Tamoxifen resistance can drive EMT



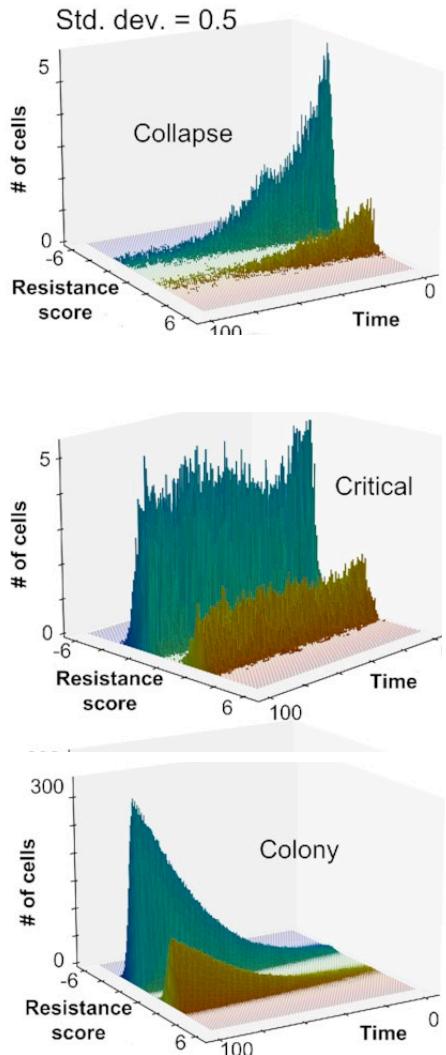
Sarthak Sahoo

Suggesting combinatorial therapies for ER+ breast cancer

A



Sahoo et al.
NAR Cancer 2021



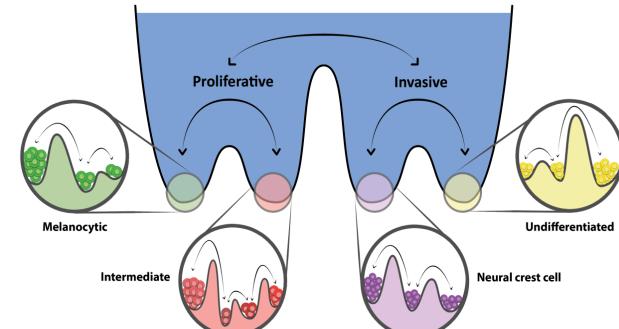
Model predictions currently undergoing experimental validation

Summary

- **Multistable** dynamics of underlying networks driving cell-state switching

⇒ Phenotypic plasticity

⇒ Non-genetic heterogeneity



Pillai & Jolly, **iScience** 2021

- ‘Design principles’ of such networks:

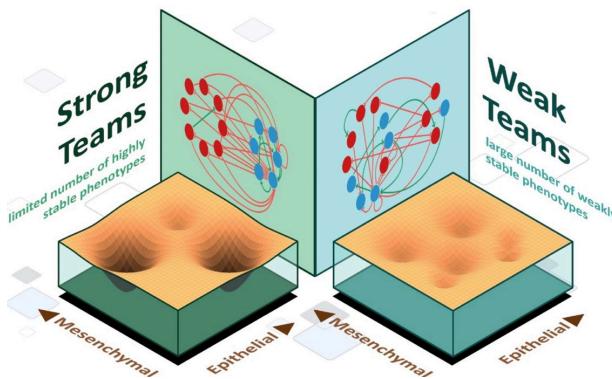
1. “Teams” exist in multiple such networks
2. “Teams” offer canalization of phenotypes



Chauhan*, Ram* et al. **eLife** 2021

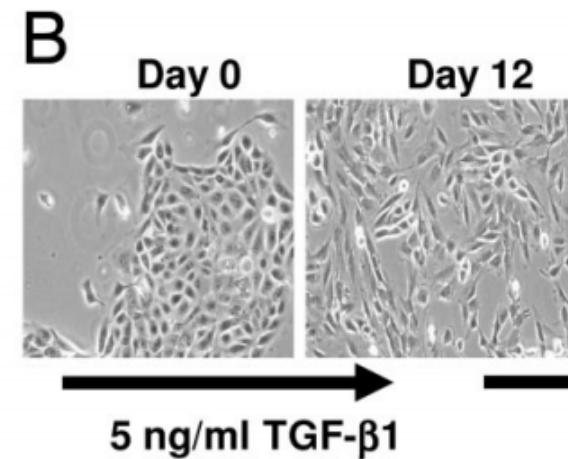
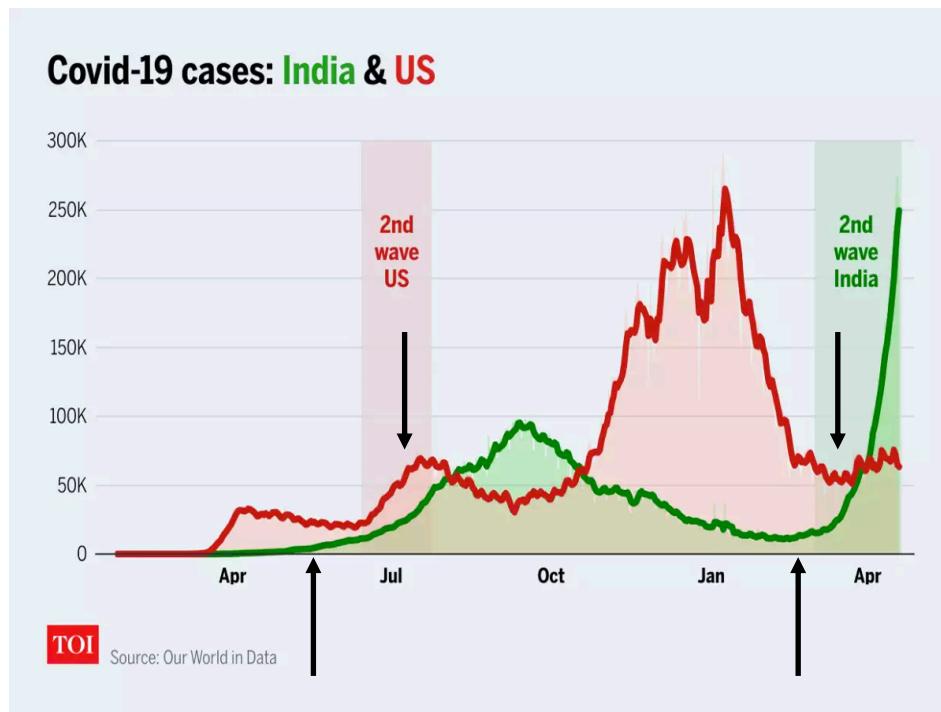
- These networks can explain **adaptive, heterogeneous response to drug treatment**

- These networks as platforms to predict combination and sequence of therapies?



Hari et al. **bioRxiv** 2021: 472090

Dynamical vs. Static hallmarks of Cancer

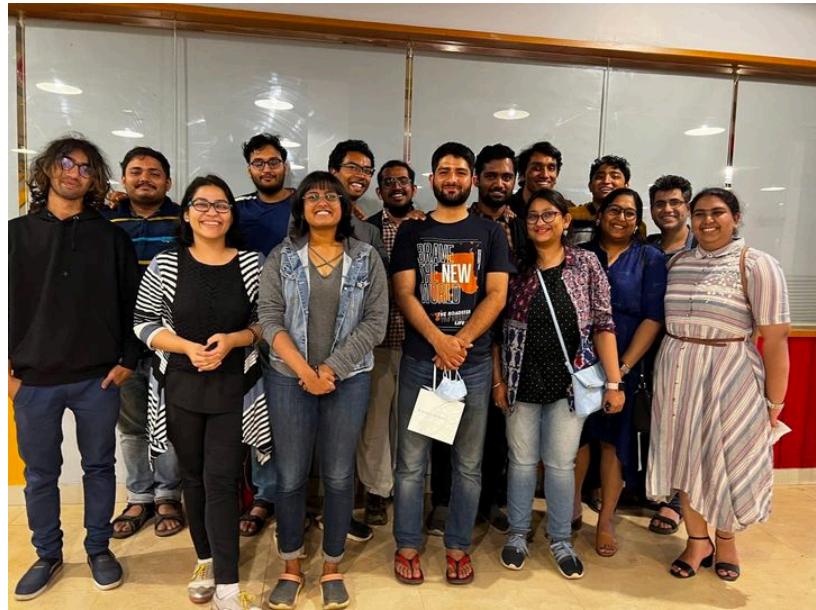


Mani et al.
PNAS 2007

Cancer is a complex, dynamic, adaptive system
(and therefore needs to be looked at such).

Math models, coupled with experimental data, have steered our understanding of such systems (weather predictions, finance etc.)

Acknowledgements: Cancer Systems Biology Lab



Any questions/comments/
suggestions welcome!

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Background of CSB members
Biotechnology, Engineering (Electrical,
Mechanical), Bioinformatics, Physics,
Mathematics, Biology, Chemistry