

LAMFA



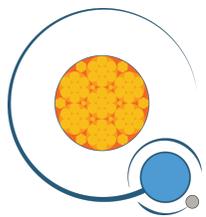
Modelling of the RAS-RAF-MEK-ERK pathway in hepatocellular carcinoma cells

Youcef Mammeri

Maître de conférences HDR

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Joint work with
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Christophe Louandre, Corinne Godin, Chloé Sauzay,
Antoine Galmiche



LAMFA

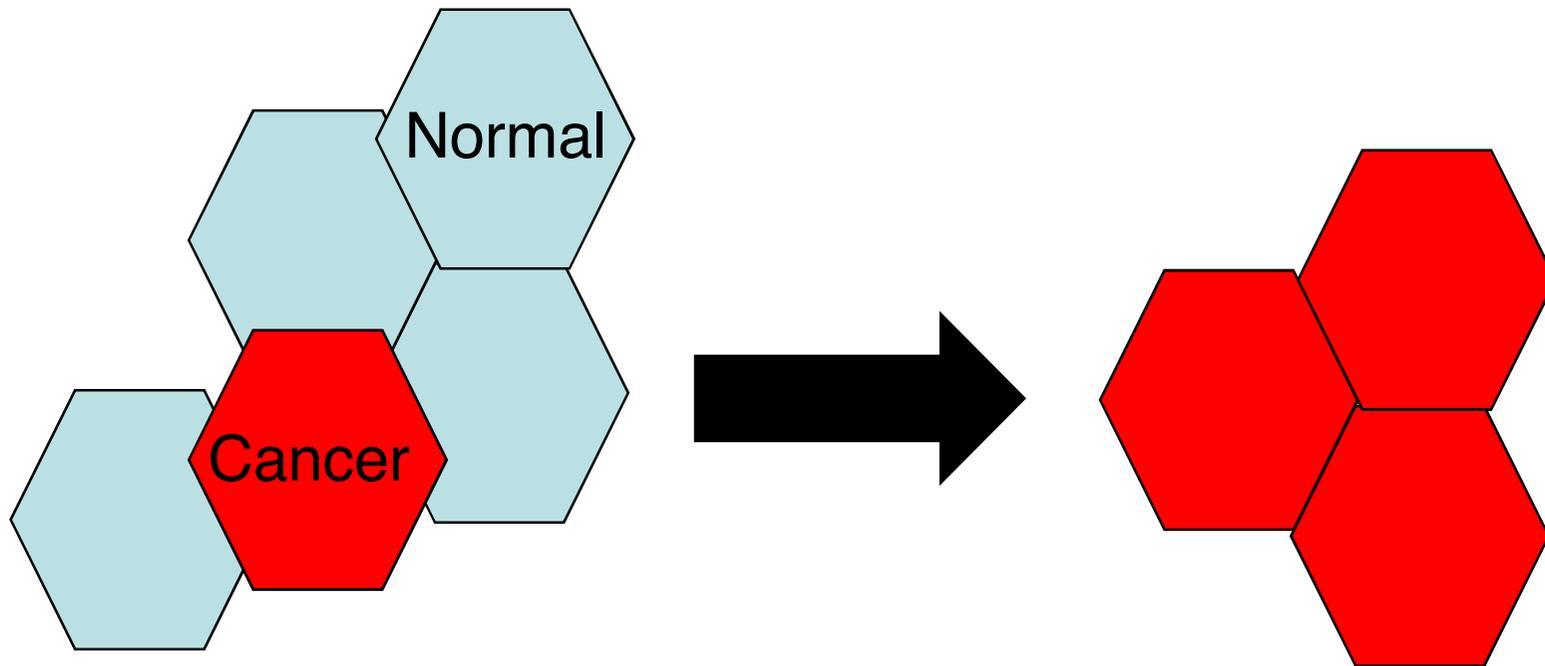


SORAMIX

Outline

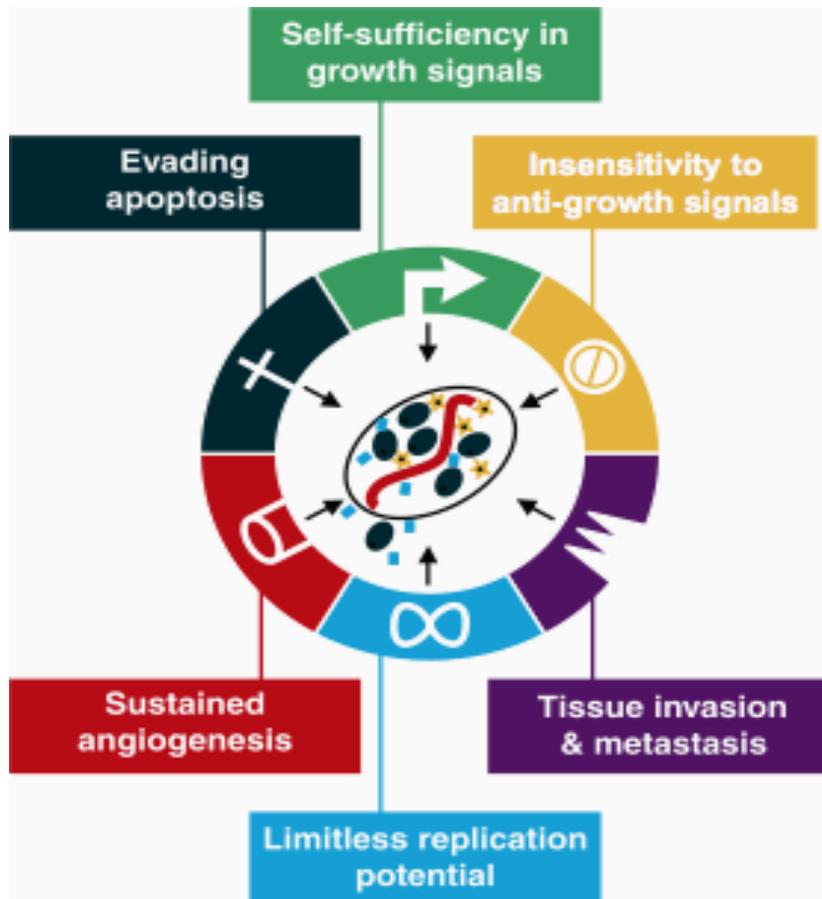
1. General informations about cancer
2. Therapeutic target: the ERK pathway
3. Modelling of the pathway
4. Conclusion

Cancer



Onset of a tumor is a clonal process +
inheritable cell phenotype

Tumor cells



Hanahan & Weinberg, Cell 2000

increased proliferation, resistance to apoptosis, invasion of tissues, ability to form metastases

Complex disease

Up to several tens of thousands of mutations in the genome of cancer cells

Each cancer is different

Sequencing compared lung cancer cell / blood cells:
23000 mutations
(10 cigarettes smoked = 1 mutation)

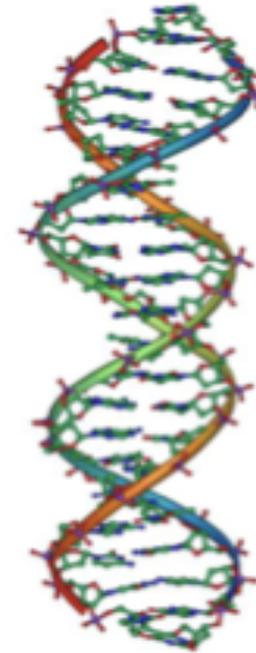


Nature 463, 2010.

An idea of the problem

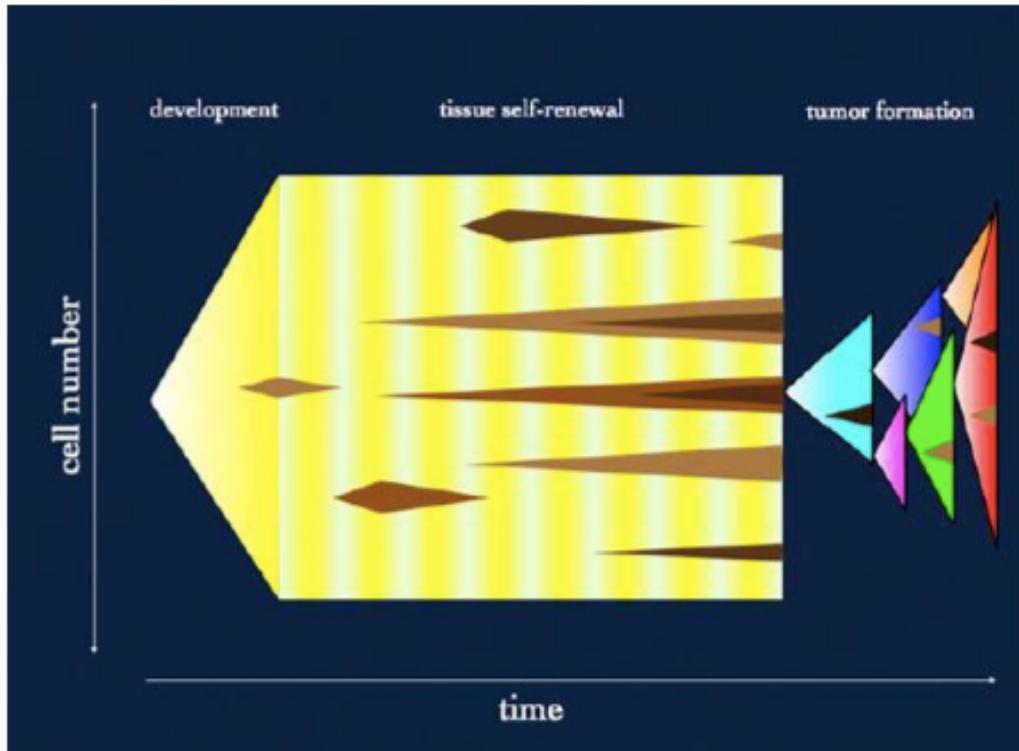


Bible:
3 million of characters



Genome:
3 billion of characters

Mutations accumulation



Proc Natl Acad Sci USA 110, 2013.

Mutations accumulation

Fraction of somatic mutations before tumor initiation

Tumor type	25-y-old	Median age at diagnosis	85-y-old
CLL	$(10.09 - 5.86)/10.09 = 0.42$	$(18.53 - 5.86)/18.53 = 0.68$	$(24.16 - 5.86)/24.16 = 0.76$
Uterine cancer	$(45.96 - 33.33)/45.96 = 0.27$	$(77.95 - 33.33)/77.95 = 0.57$	$(96.48 - 33.33)/96.48 = 0.65$
Colorectal cancer	$(50.2 - 50.2)/50.2 = 0$	$(102.4 - 50.2)/102.4 = 0.51$	$(121.38 - 50.2)/121.38 = 0.59$

time

Proc Natl Acad Sci USA 110, 2013.

Mutations accumulation

Fraction of somatic mutations before tumor initiation

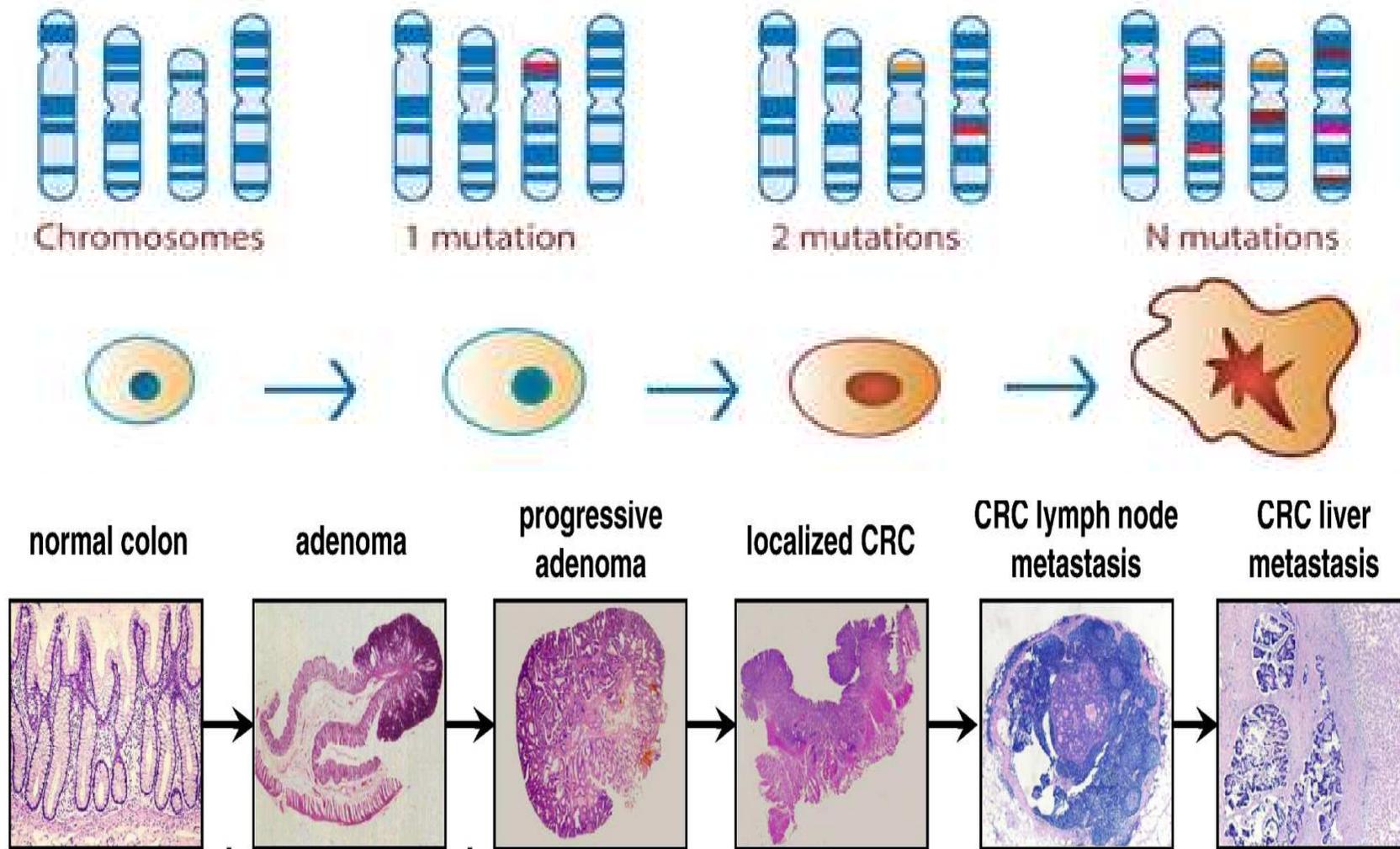
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time

Proc Natl Acad Sci USA 110, 2013.

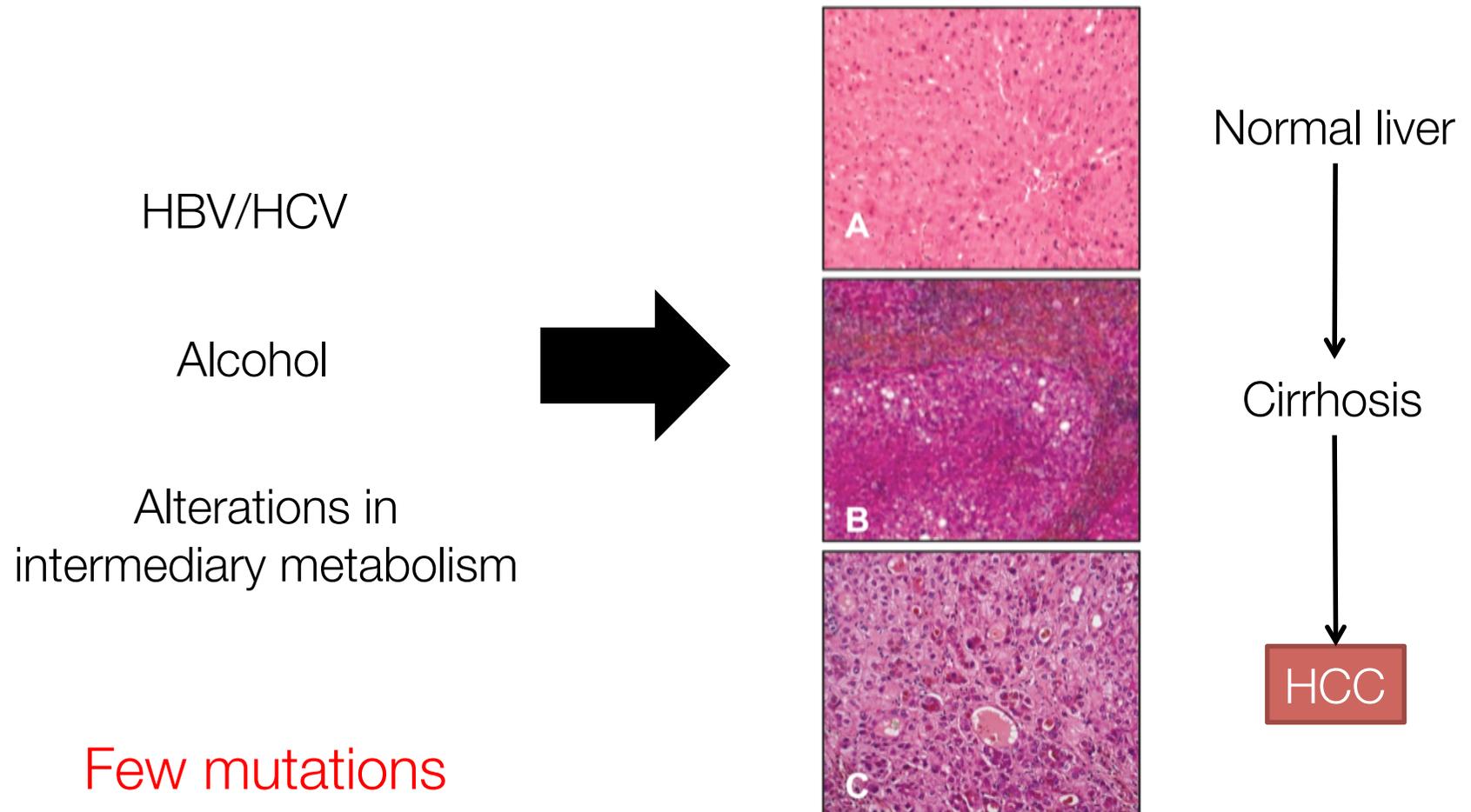
Most mutations precede the onset of cancer!

Mutations accumulation



Hepatocellular carcinoma cells

HCC the most common form of primary liver tumor



Outline

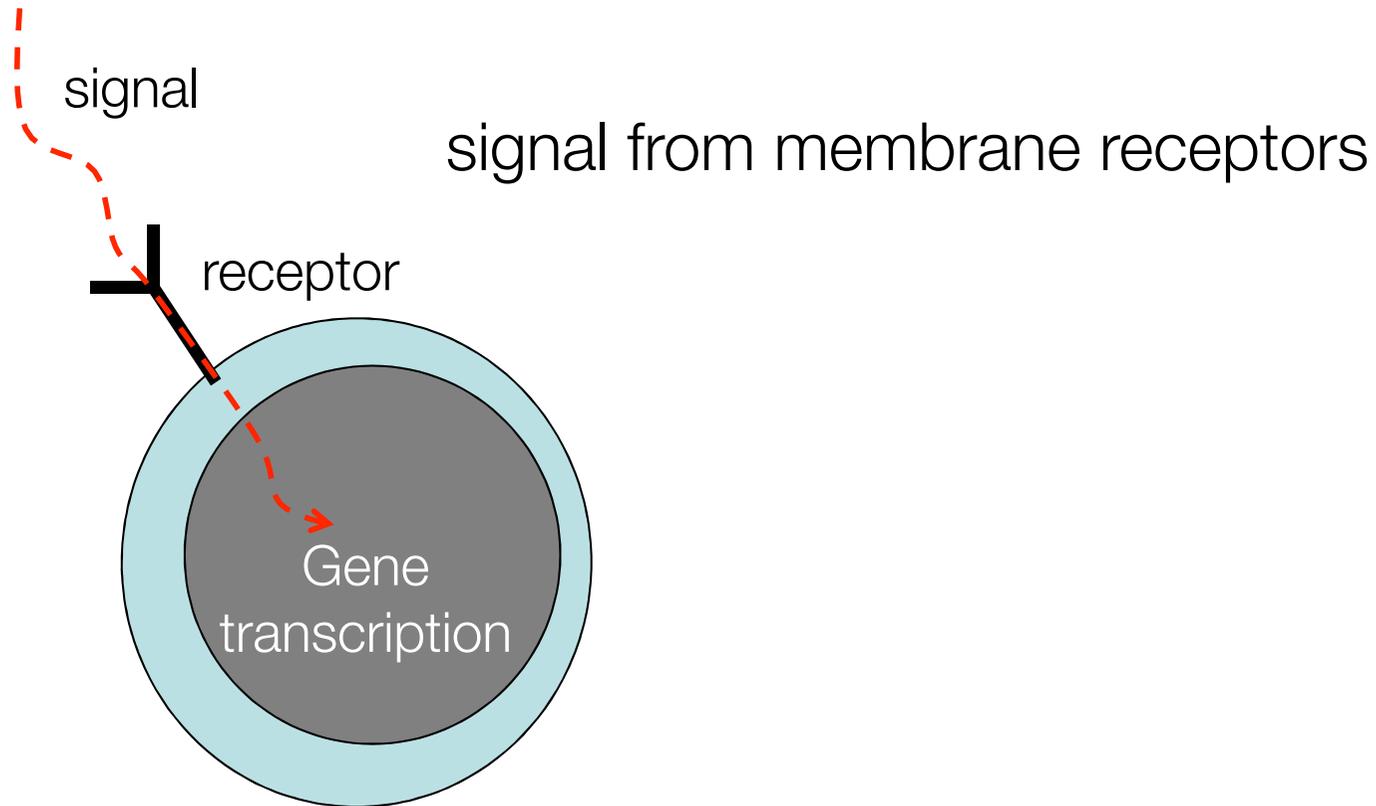
1. General informations about cancer

2. Therapeutic target: the ERK pathway

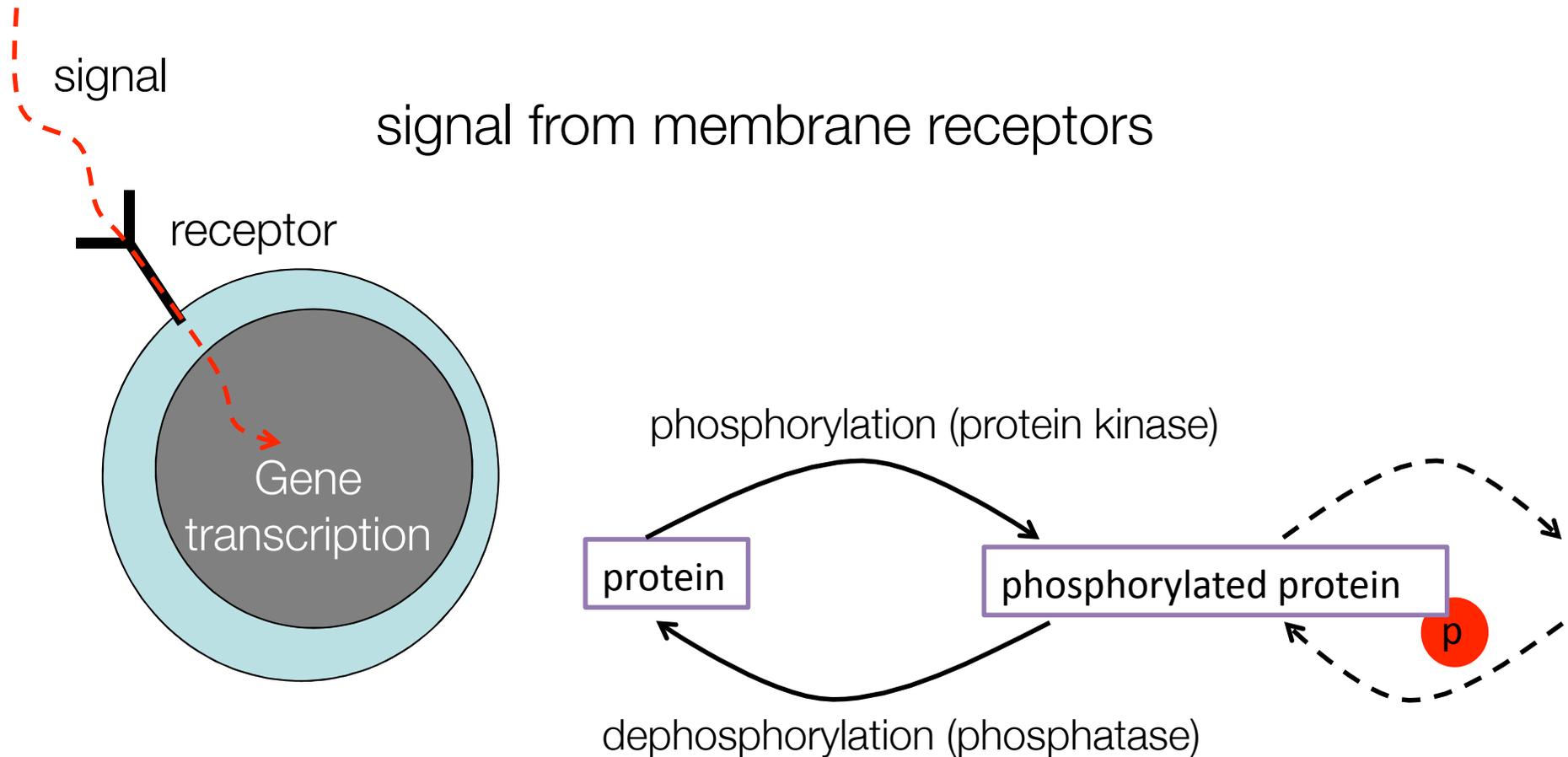
3. Modelling of the pathway

4. Conclusion

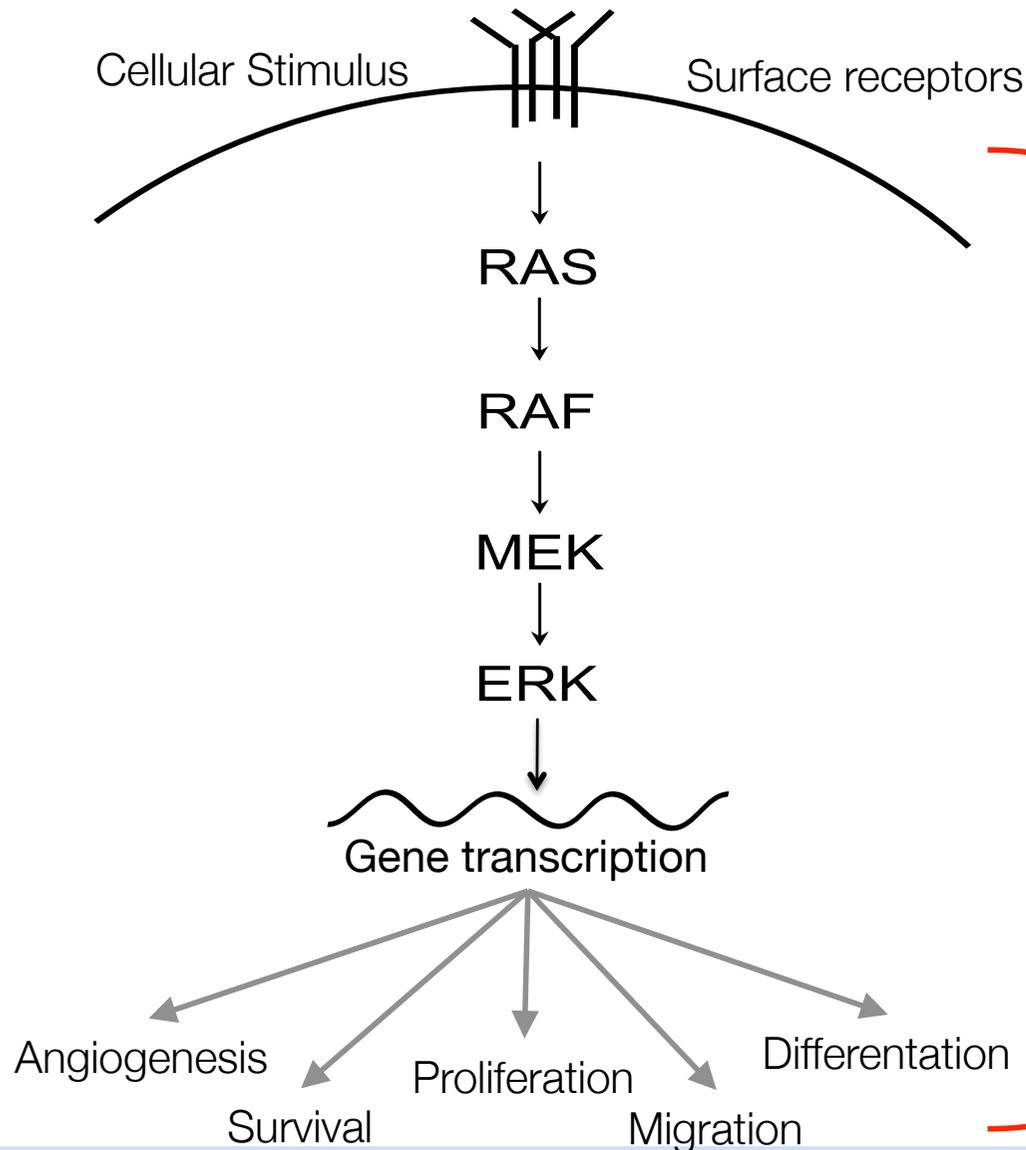
Cell signaling



Cell signaling

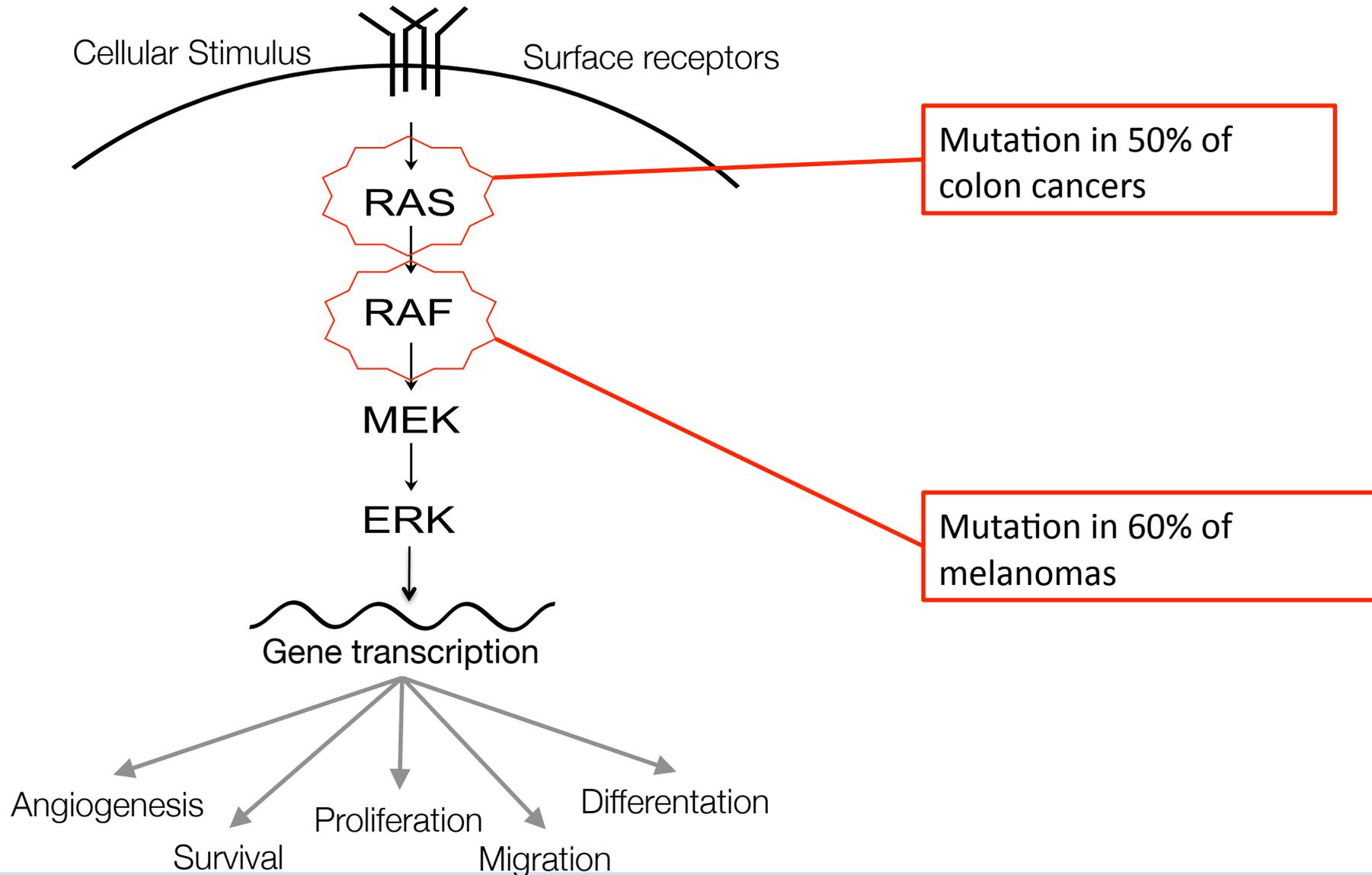


Basic pathway



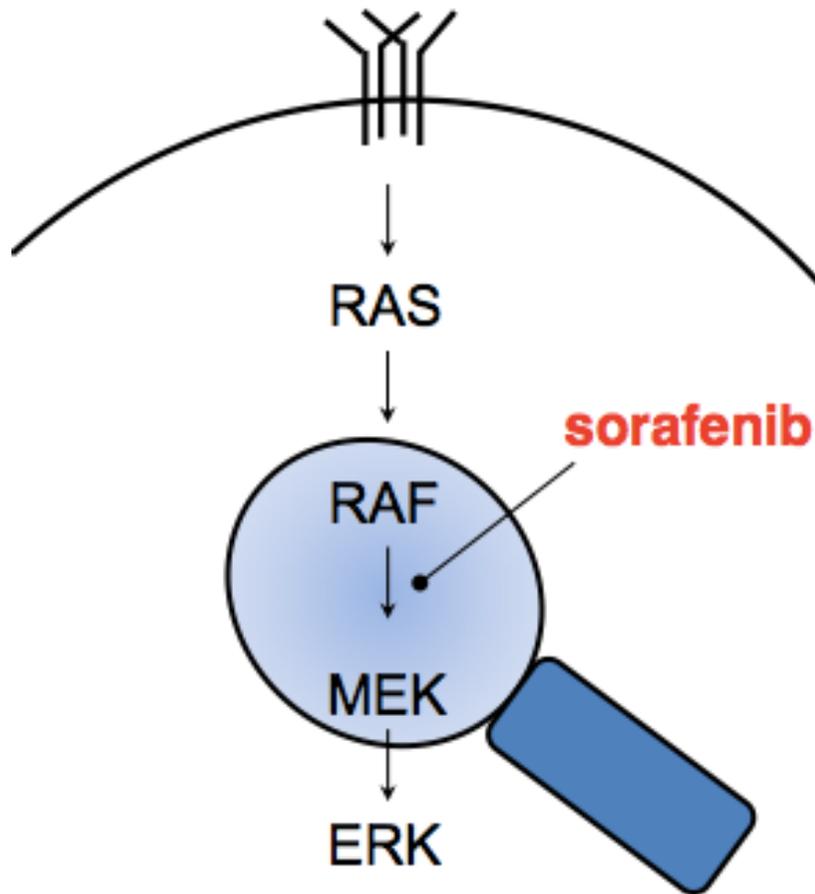
A normal but often deviated transduction pathway in cancers

Basic pathway

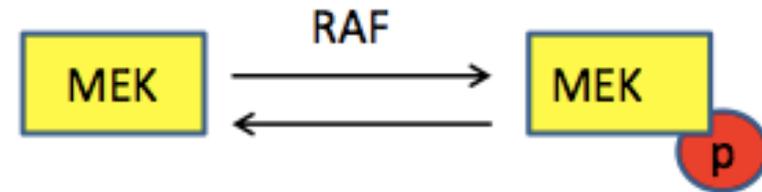


Therapeutic targeting

New medical treatments designed rationally:



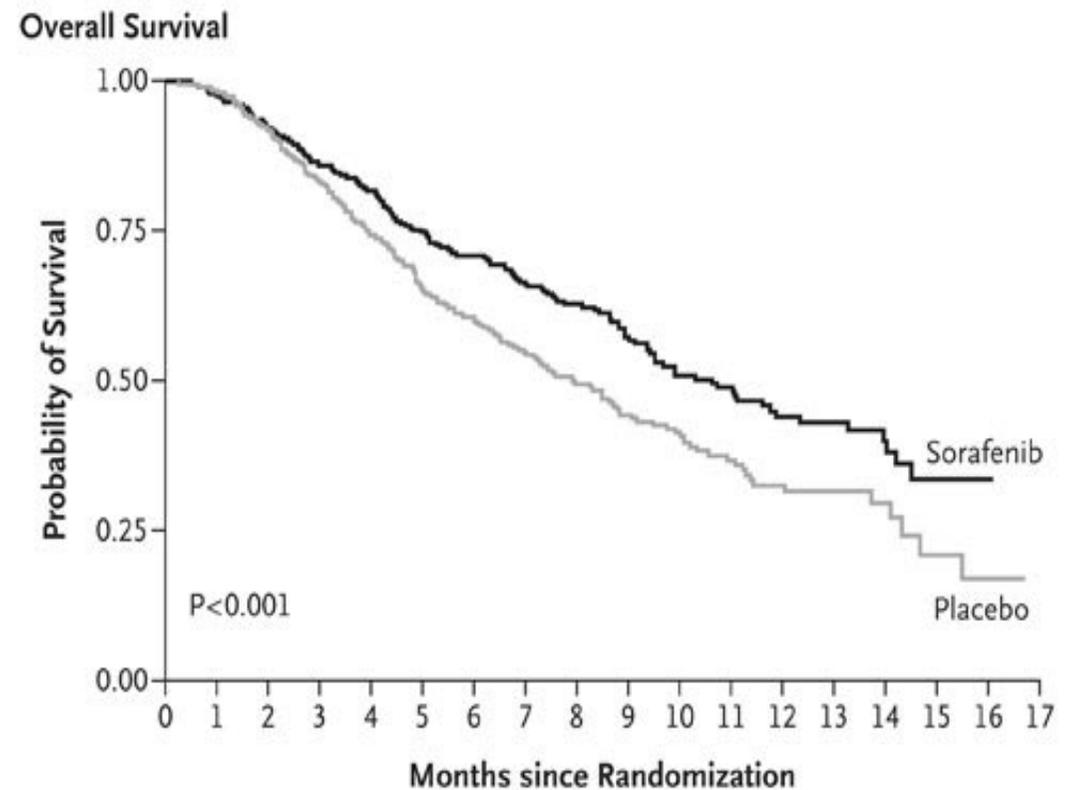
Enzyme inhibitors (kinases)



Goal: turn off oncogenic signaling

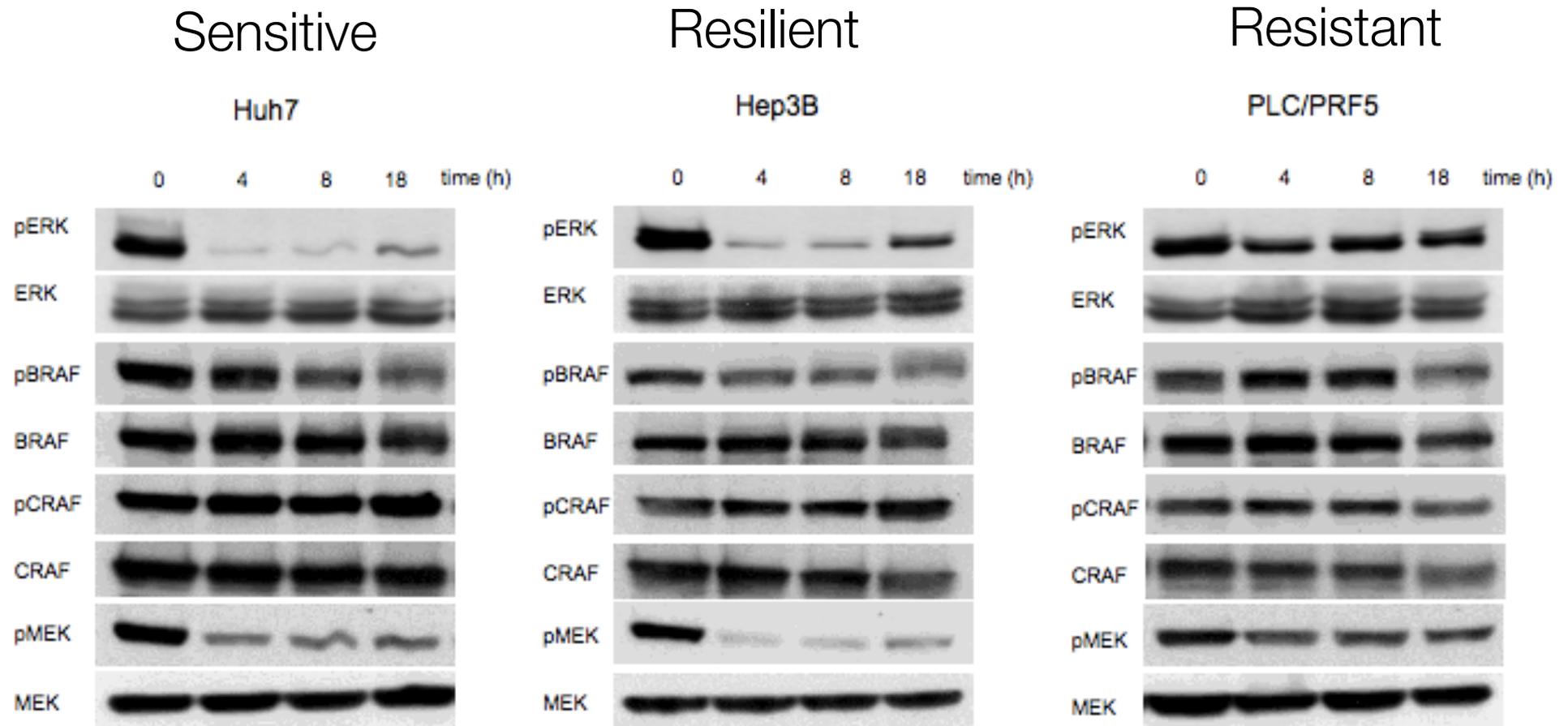
Sorafenib in HCC

- Mortality +++
- 80% of patients are diagnosed at a late stage of the disease
- Until 2008, no medical treatment of proven efficacy on survival for these advanced stages



Llovet et al. N Engl J Med 2008

Heterogeneous dynamic



Saidak et al. Cancer Letters 2017

Motivation

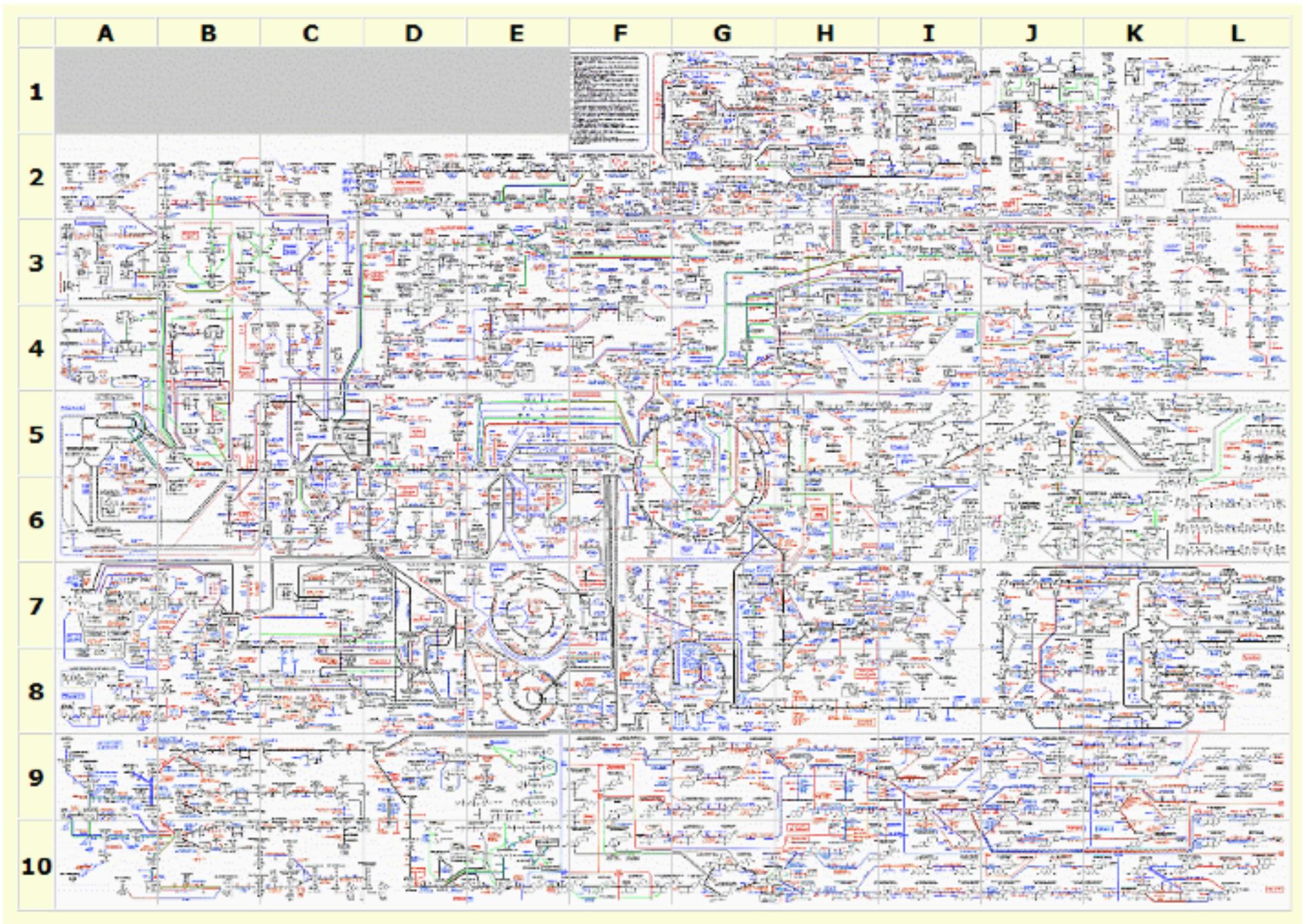
Improve the knowledge of the ERK pathway regulation

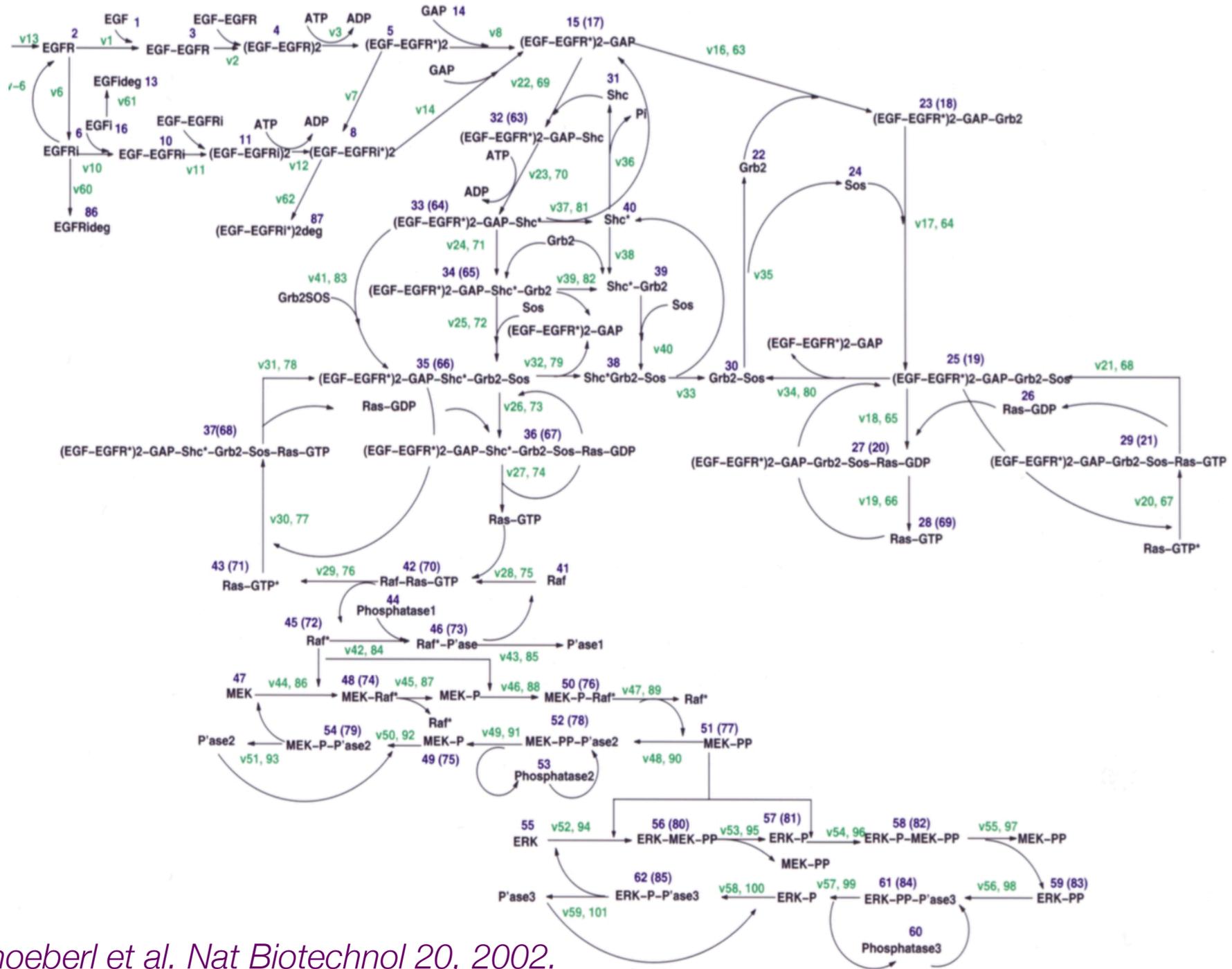
Understanding how the medicine works

Identify and control the pathway

Outline

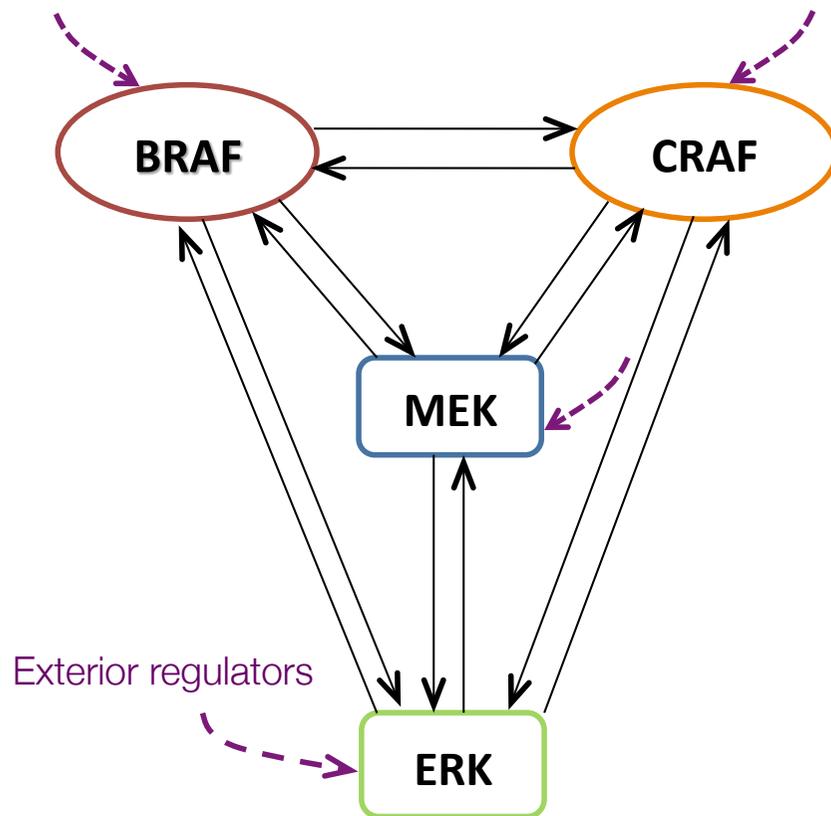
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Schoeberl et al. Nat Biotechnol 20, 2002.

Average



Material support

3 cell lines : Hep3b (sensitive), Huh7 (resilient) and PLC\PRF5 (resistant),

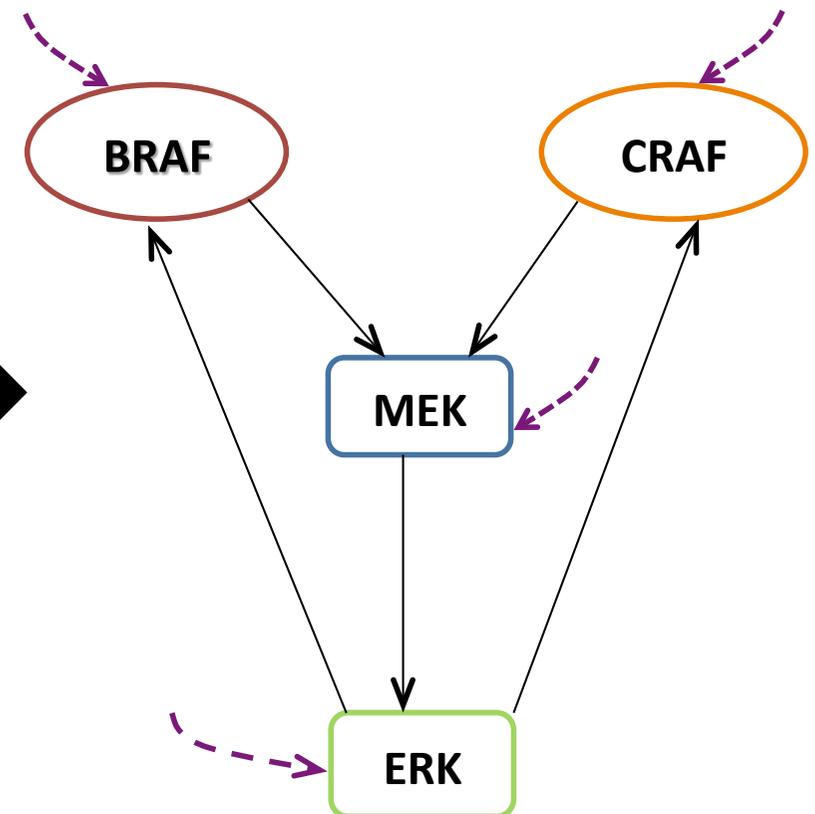
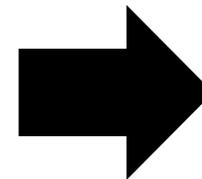
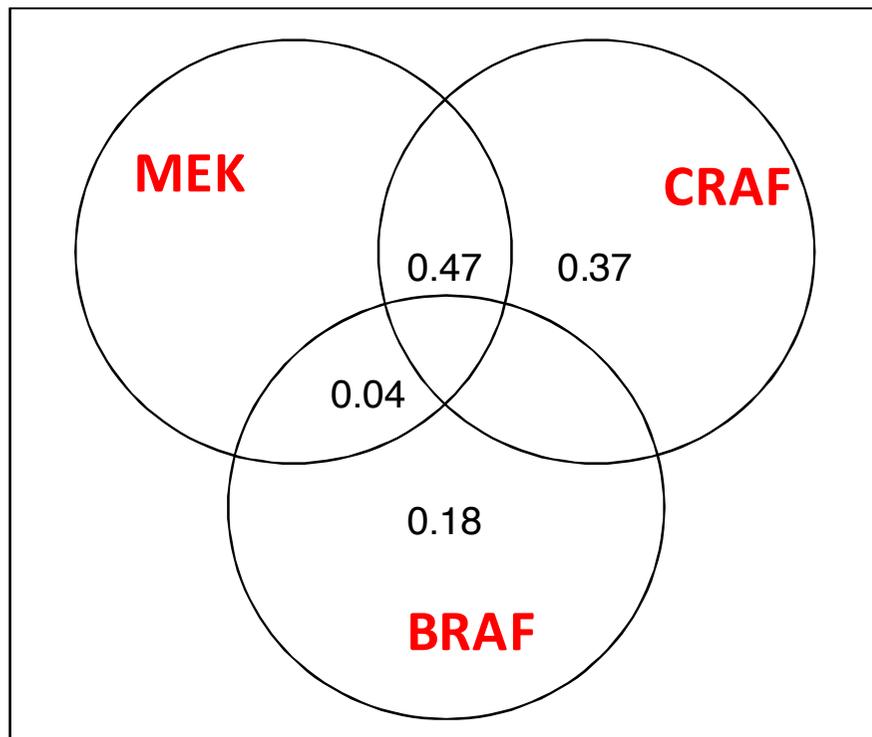
Western blot analysis

Quantification of each phosphorylated protein and their total at three measure times after sorafenib treatment (4, 8 and 18h).

Experiments repeated three times.

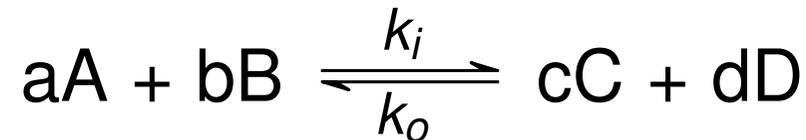
Data analysis

ERK at 18 hours



Model a reaction

The general chemical reaction



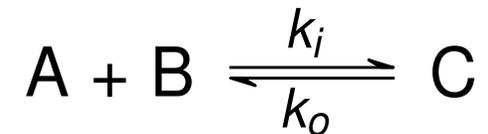
defines a reaction rate

$$v = -\frac{1}{a} \frac{d[A]}{dt} = -\frac{1}{b} \frac{d[B]}{dt} = \frac{1}{c} \frac{d[C]}{dt} = \frac{1}{d} \frac{d[D]}{dt}$$

where $[X]$ denotes the concentration of the chemical species X (usually in mol/L).

Model a reaction

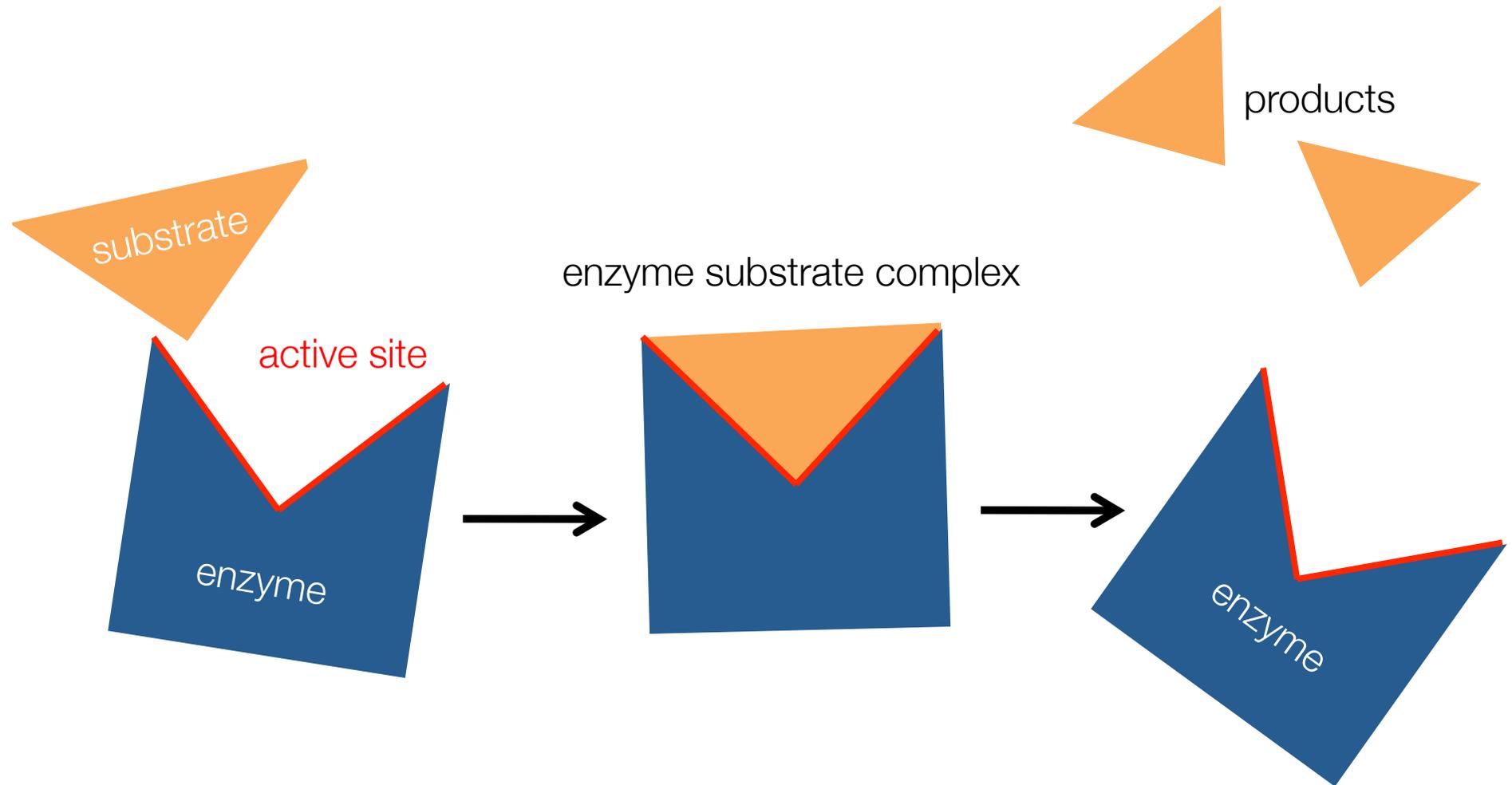
Consider the simple reaction



the change in amount of [A] = production – degradation

$$\frac{d[A]}{dt} = \frac{d[B]}{dt} = -\frac{d[C]}{dt} = k_o[C] - k_i[A][B]$$

Enzyme kinetics



Enzyme kinetics



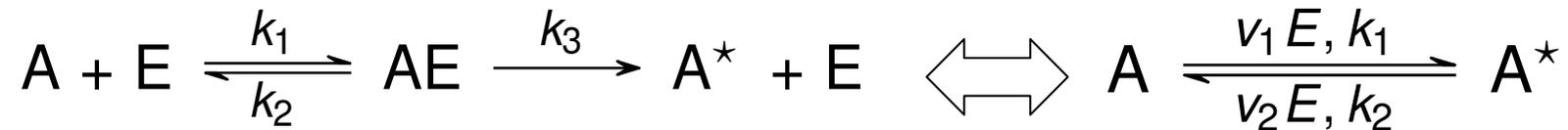
$$\frac{d[A]}{dt} = -k_1[A][E] + k_2[AE]$$

$$\frac{d[E]}{dt} = -k_1[A][E] + k_2[AE] + k_3[AE]$$

$$\frac{d[AE]}{dt} = k_1[A][E] - k_2[AE] - k_3[AE]$$

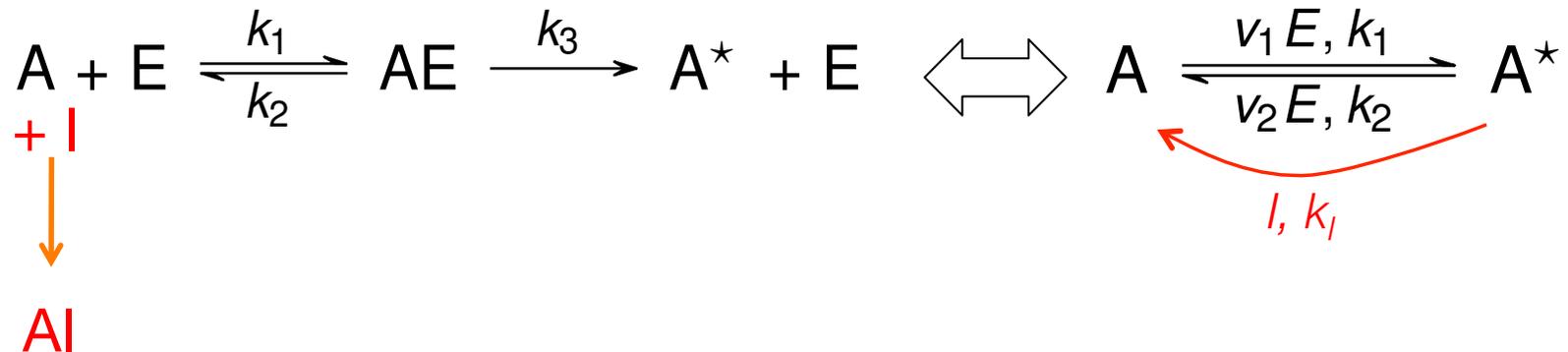
$$\frac{d[A^*]}{dt} = k_3[AE]$$

Enzyme kinetics



$$\frac{d[A^*]}{dt} = \frac{v_1 [E] ([A_{total}] - [A^*])}{k_1 + ([A_{total}] - [A^*])} - \frac{v_2 [A^*]}{k_2 + [A^*]}$$
$$A_{total} = A + A^*.$$

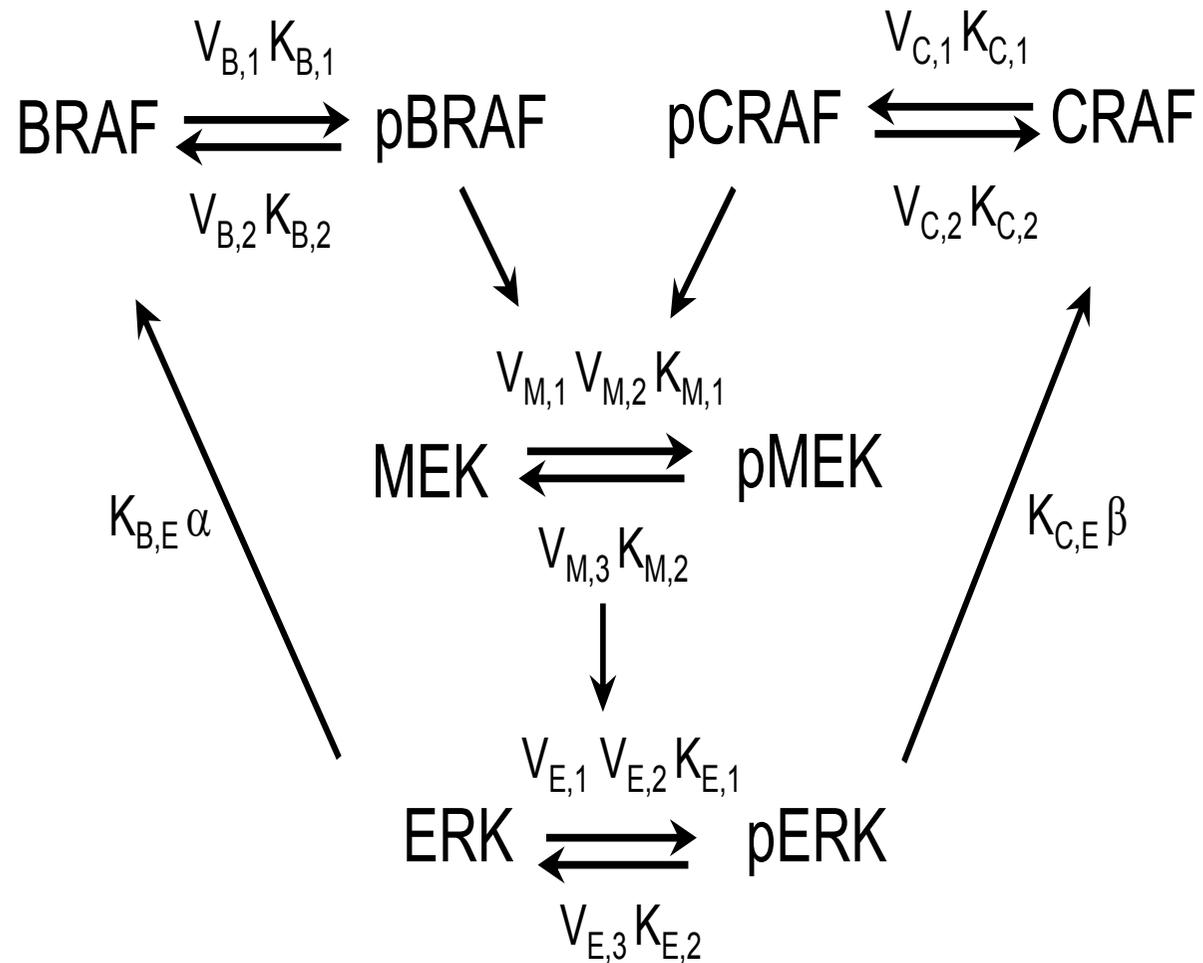
Enzyme kinetics



$$\frac{d[A^*]}{dt} = \frac{v_1 [E] ([A_{total}] - [A^*])}{k_1 + ([A_{total}] - [A^*])} \left(1 + \frac{[I]}{k_I} \right)^{\alpha_I} - \frac{v_2 [A^*]}{k_2 + [A^*]}$$

$$A_{total} = A + A^* .$$

ERK pathway



ERK pathway

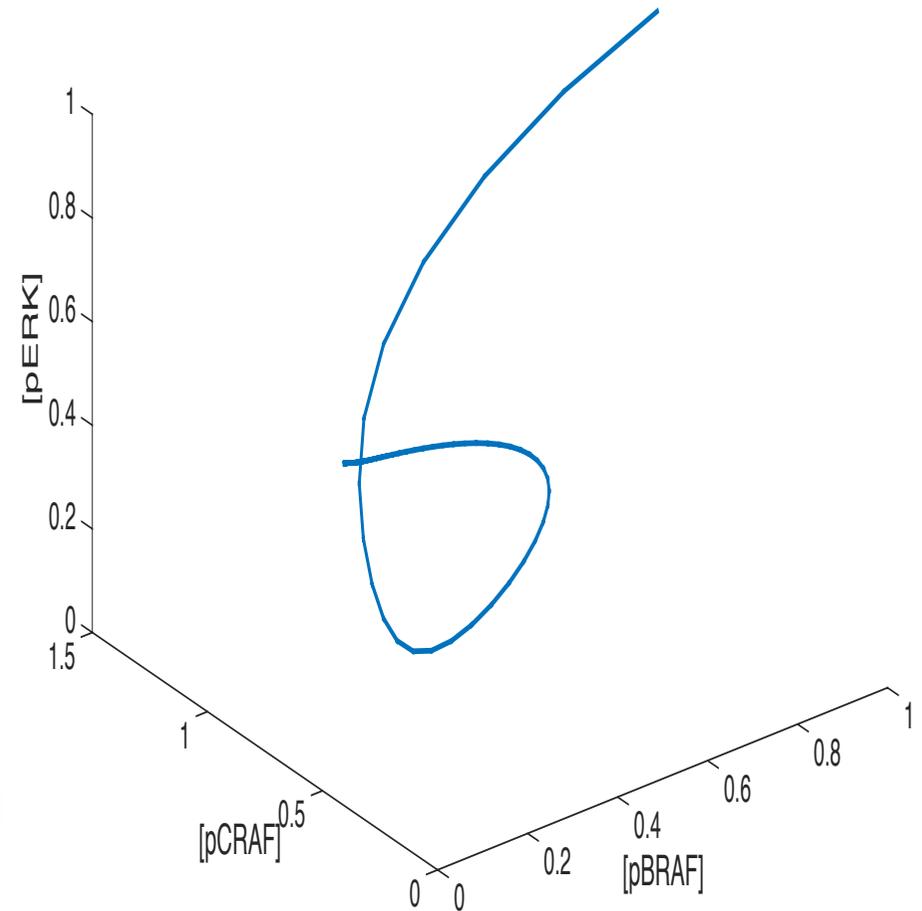
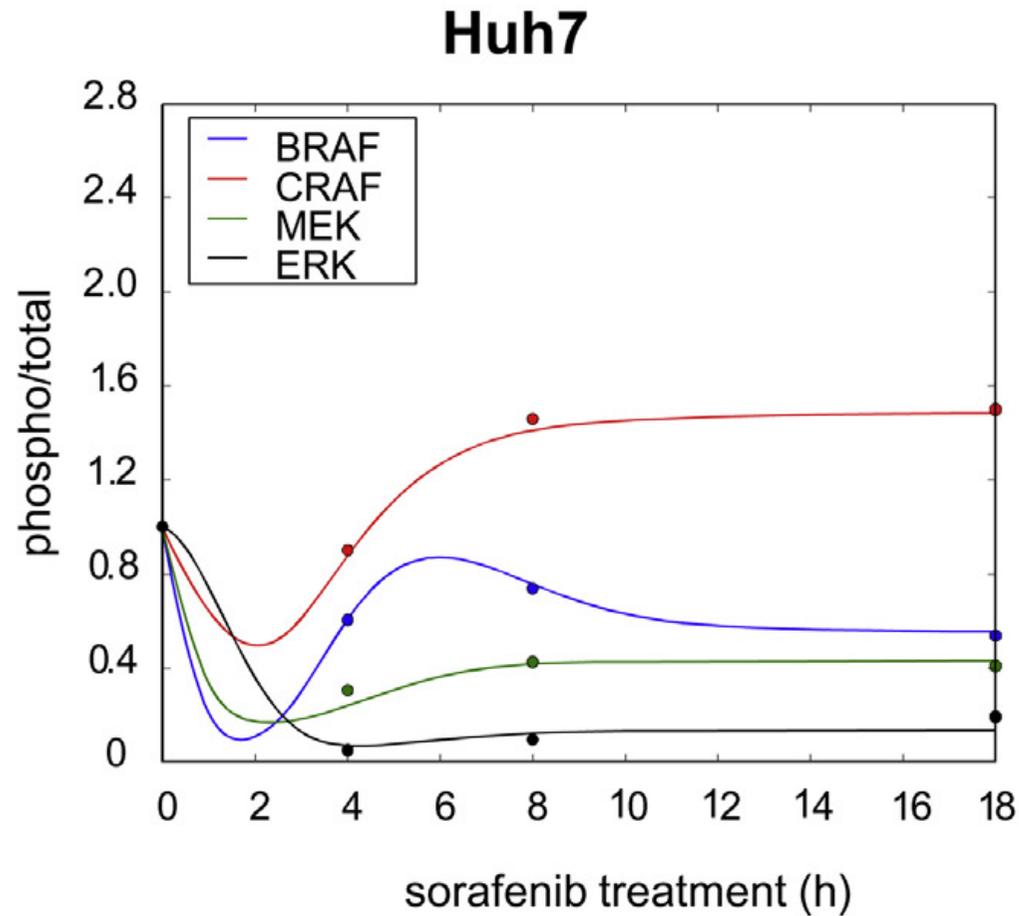
$$\frac{d[pBRAF]}{dt} = \frac{V_{B,1} ([BRAF_{tot}] - [pBRAF])}{(K_{B,1} + ([BRAF_{tot}] - [pBRAF]))} \left(1 + \frac{[pERK]}{K_{B,E}}\right)^{-\alpha} - \frac{V_{B,2} [pBRAF]}{(K_{B,2} + [pBRAF])}$$

$$\frac{d[pCRAF]}{dt} = \frac{V_{C,1} ([CRAF_{tot}] - [pCRAF])}{(K_{C,1} + ([CRAF_{tot}] - [pCRAF]))} \left(1 + \frac{[pERK]}{K_{C,E}}\right)^{-\beta} - \frac{V_{C,2} [pCRAF]}{(K_{C,2} + [pCRAF])}$$

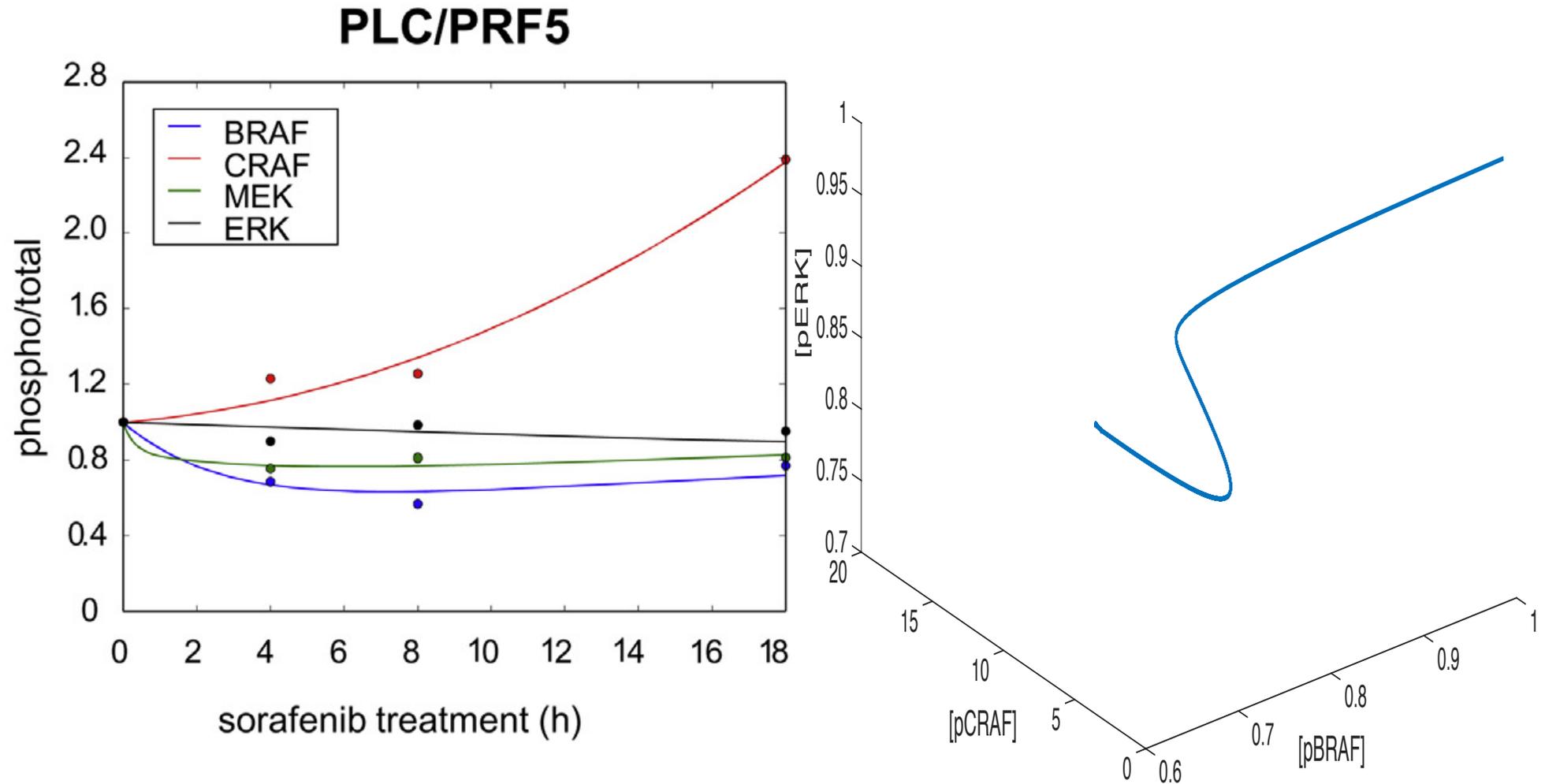
$$\frac{d[pMEK]}{dt} = \frac{(V_{M,1} + V_{M,2} [pBRAF] + (1 - V_{M,2}) [pCRAF]) ([MEK_{tot}] - [pMEK])}{K_{M,1} + ([MEK_{tot}] - [pMEK])} - \frac{V_{M,3} [pMEK]}{(K_{M,2} + [pCRAF])}$$

$$\frac{d[pERK]}{dt} = \frac{(V_{E,1} + V_{E,2} [pMEK]) ([ERK_{tot}] - [pERK])}{K_{E,1} + ([ERK_{tot}] - [pERK])} - \frac{V_{E,3} [pERK]}{(K_{E,2} + [pERK])}$$

Sensitive cells

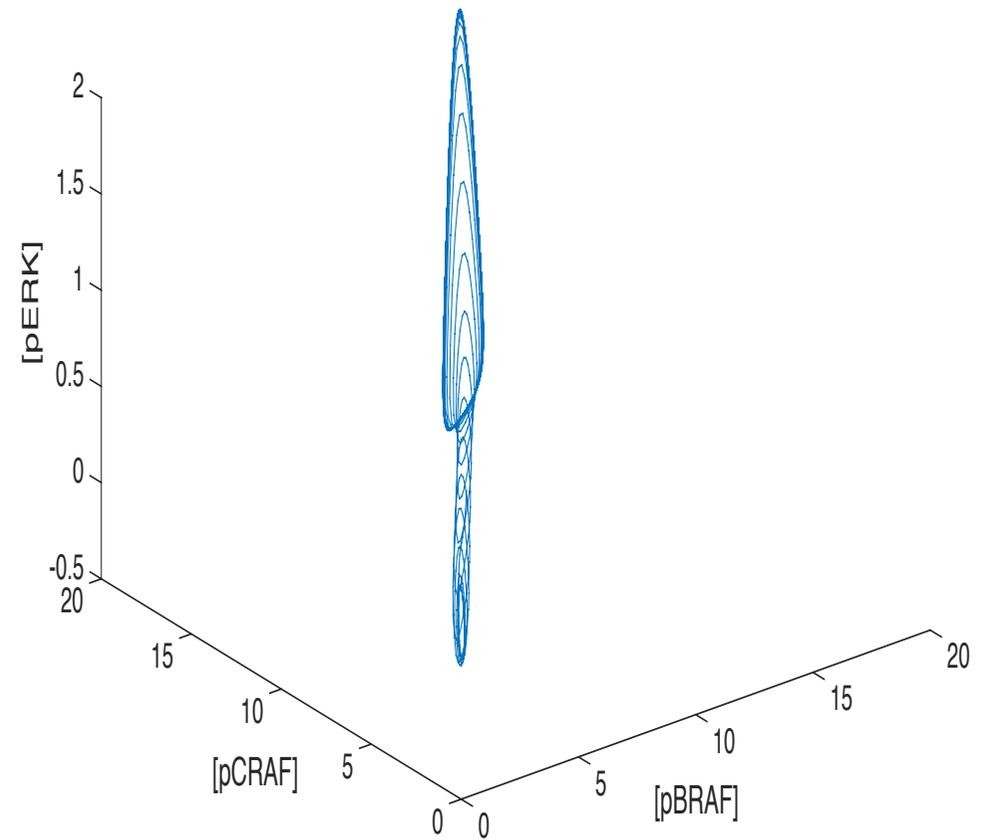
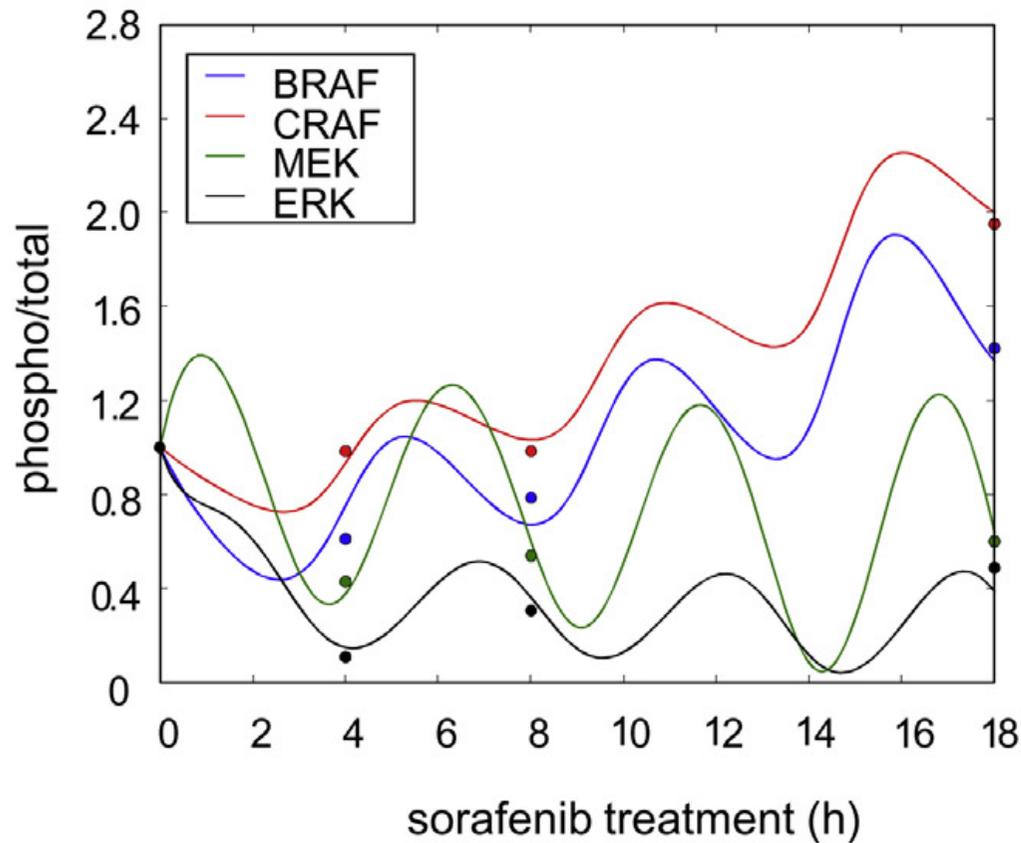


Resistant cells



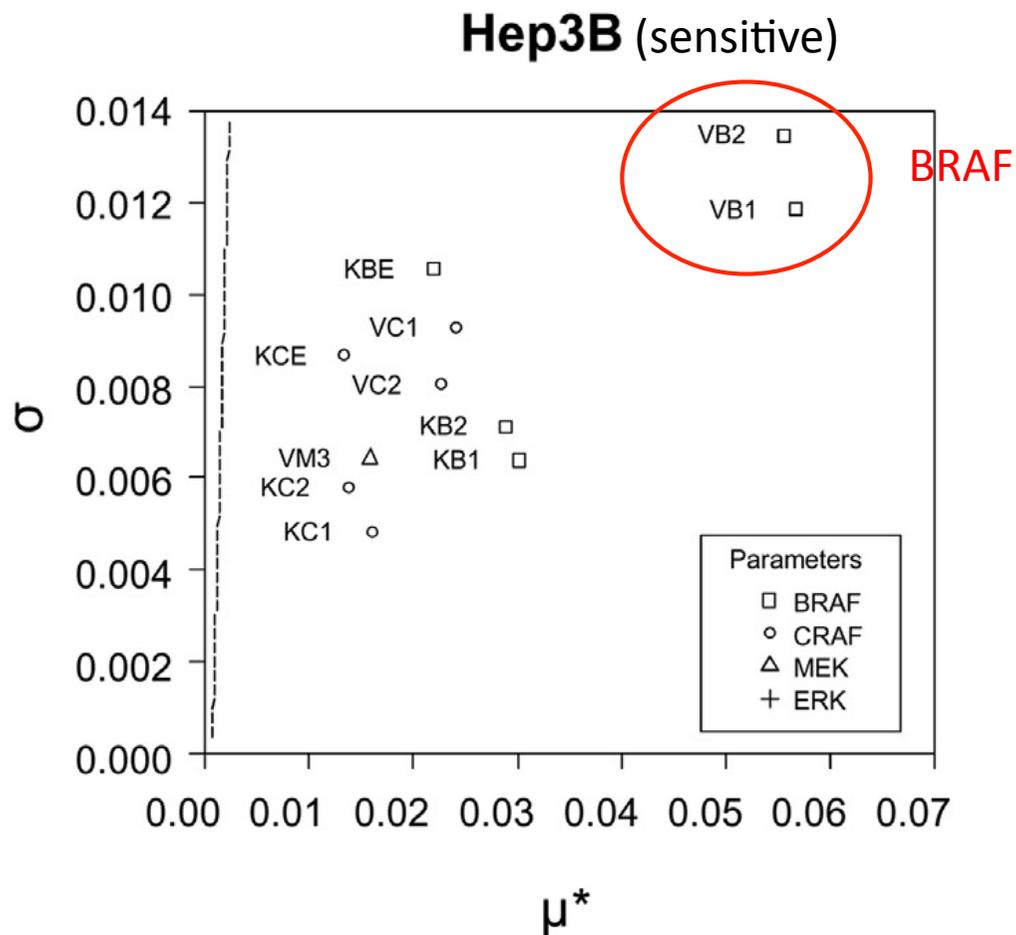
Resilient cells

Hep3B



Sensitivity analysis

Identify the most significant effects provided by the parameters on the phosphorylated ERK amount.

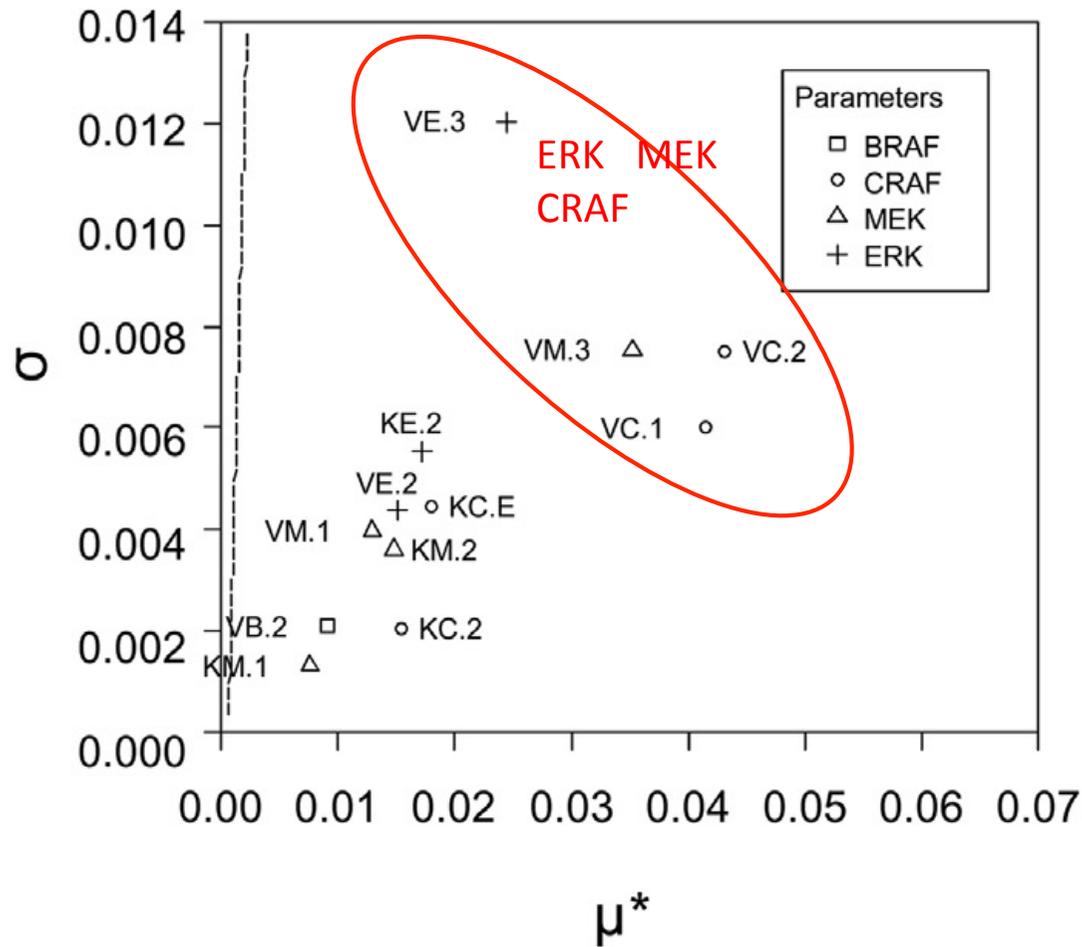


μ^* absolute mean tendancy

σ nonlinear effects with interactions

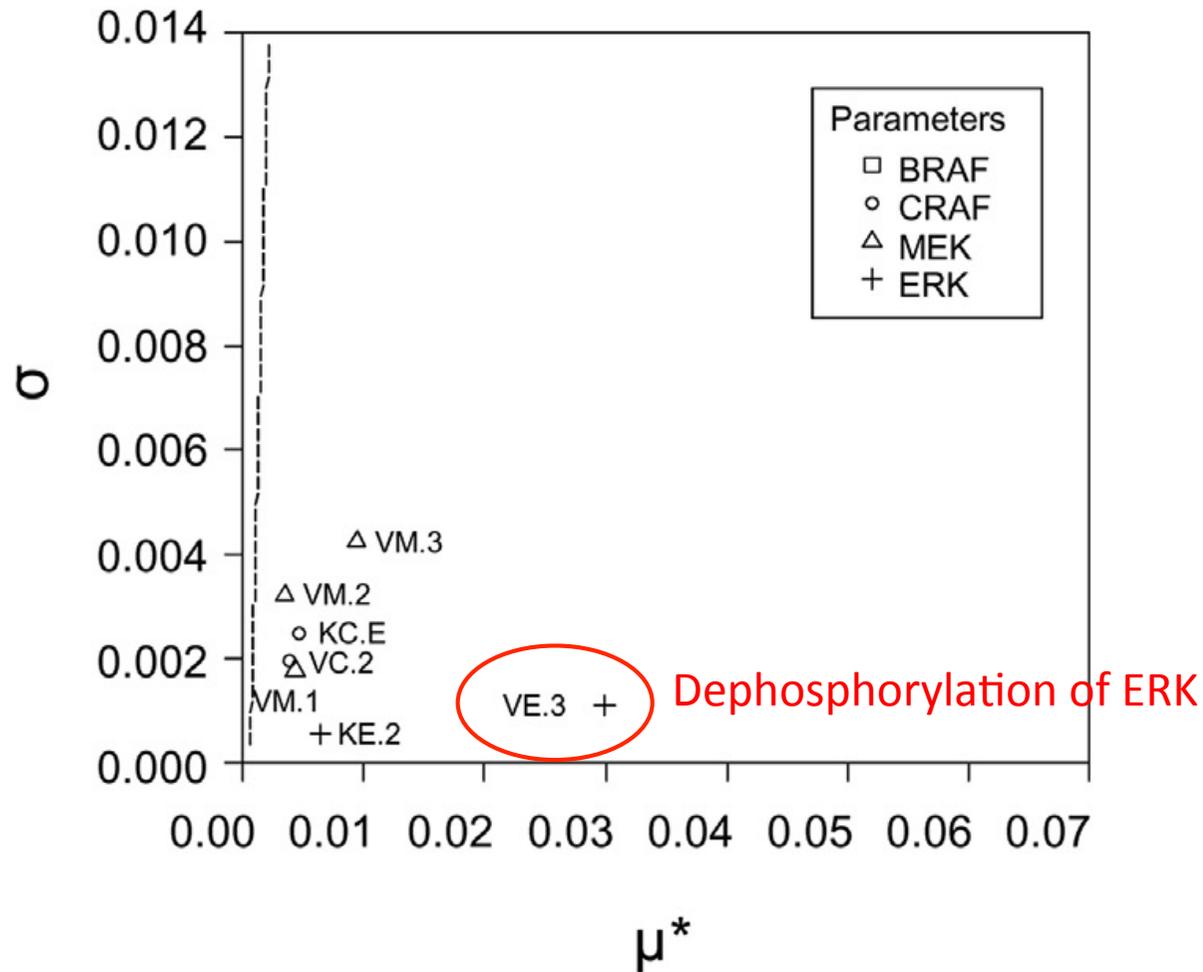
Sensitivity analysis

Huh7 (resilient)

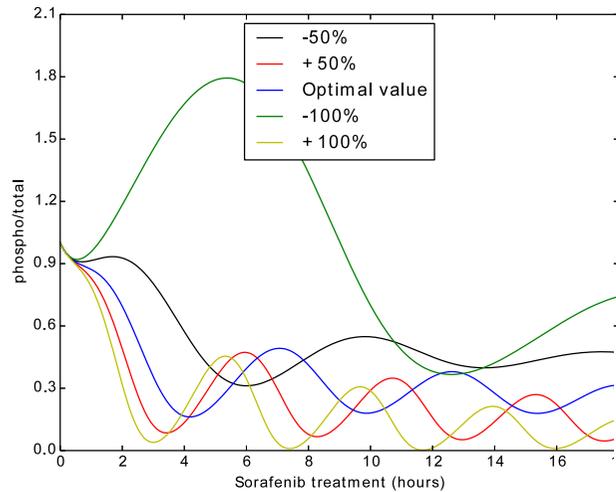


Sensitivity analysis

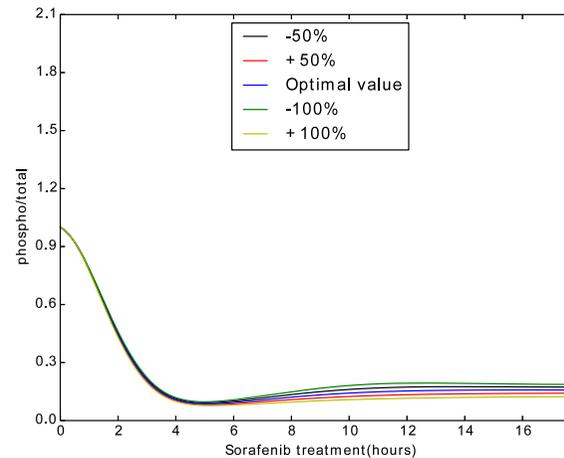
PLC/PRF5 (resistant)



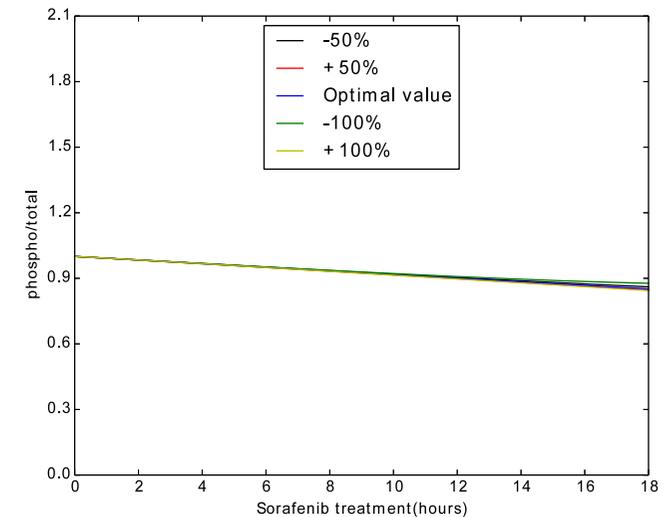
Target: RAF production?



Hep3b

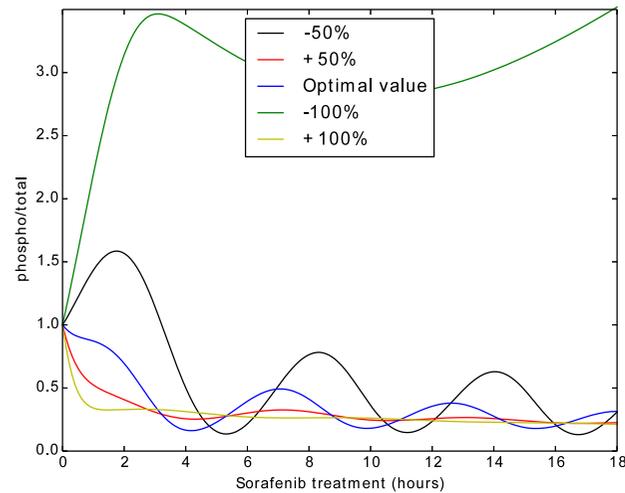


Huh7

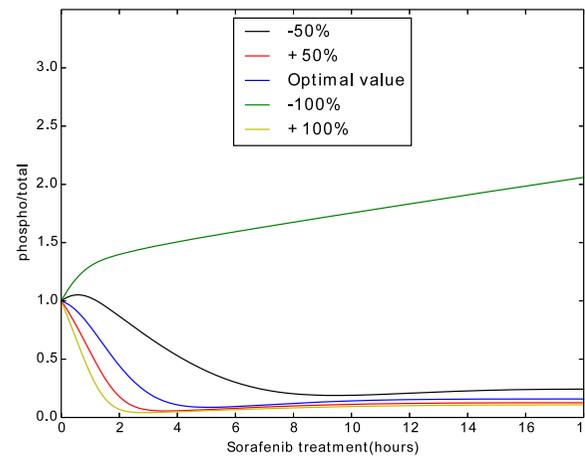


PLC\PRF5

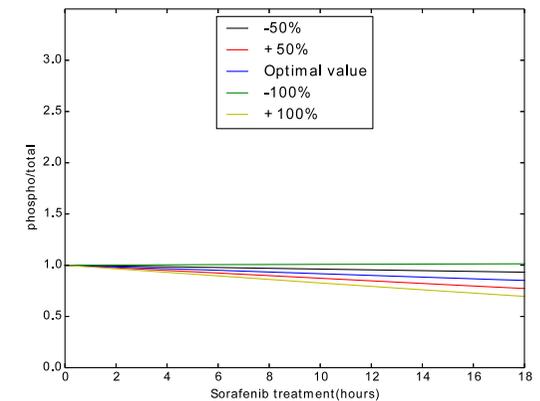
Target: ERK dephosphorylation?



Hep3b



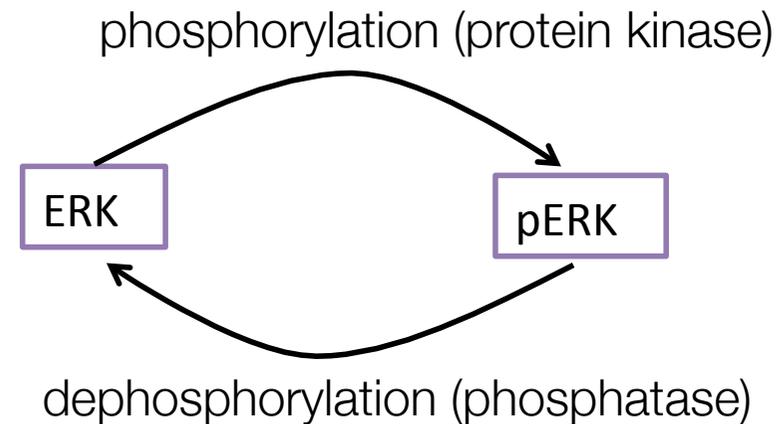
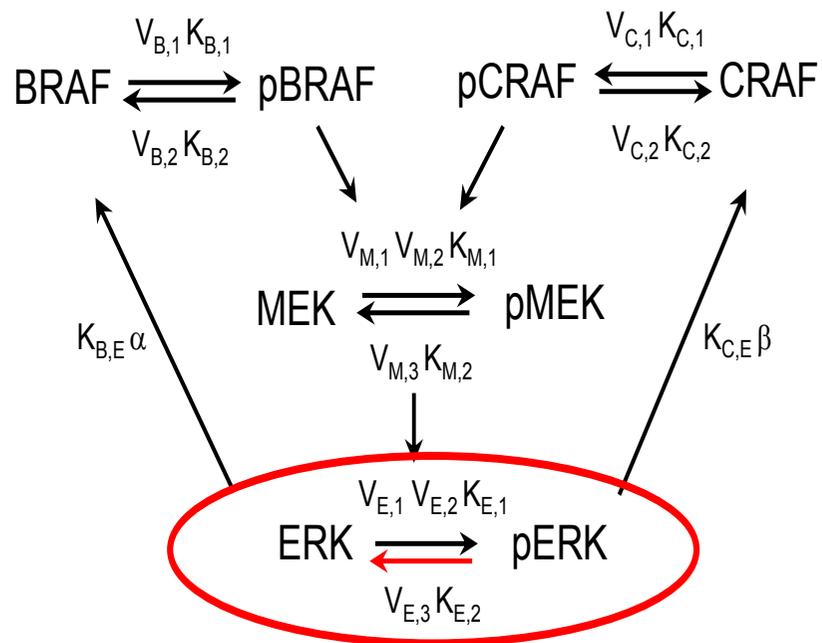
Huh7



PLC\PRF5

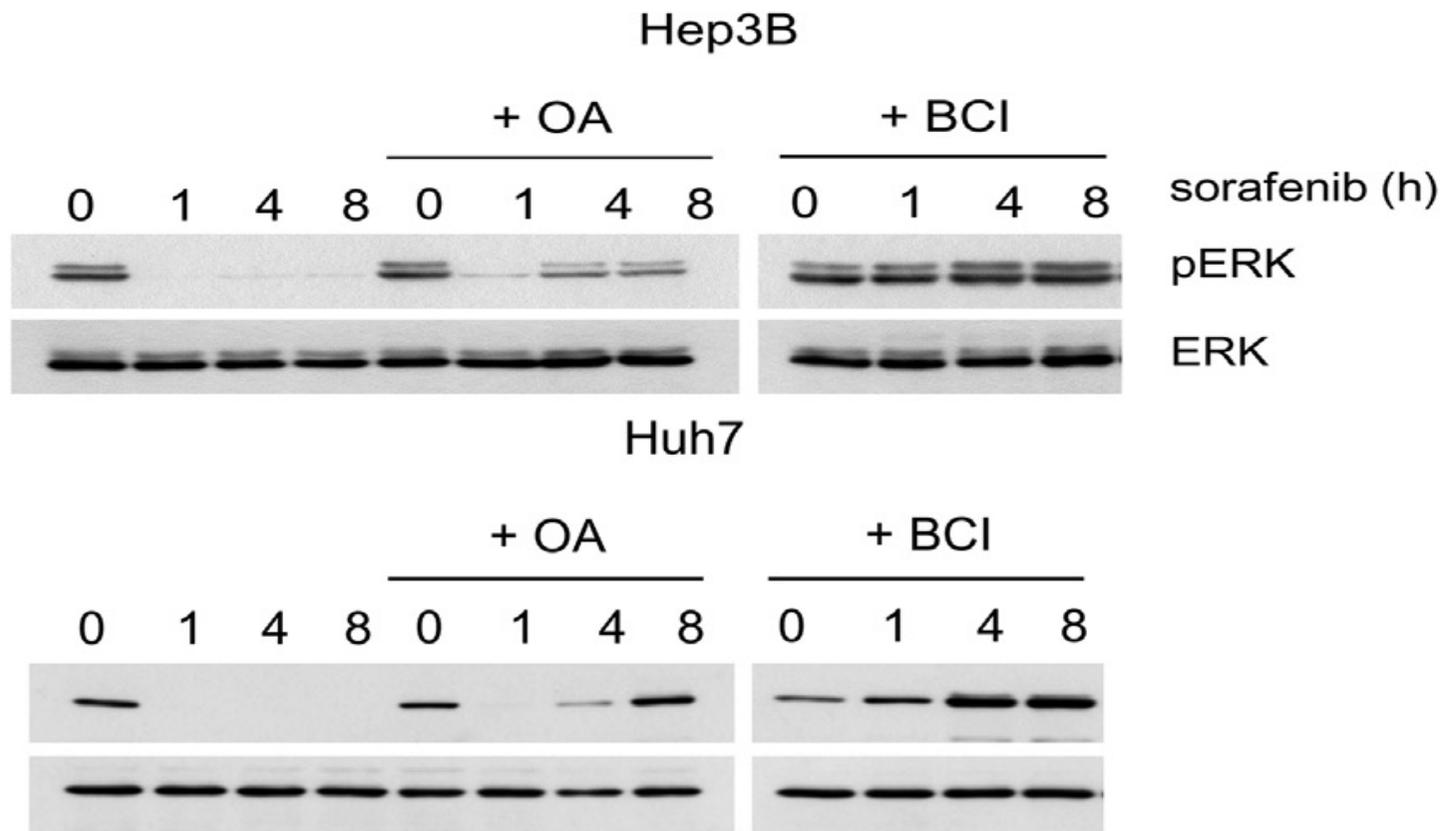
Phosphatases

An essential actor in blocking ERK pathway by sorafenib!



Phosphatases

An essential actor in blocking ERK pathway by sorafenib!



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Conclusion

Mathematics allows the study of all aspects of oncology from the most practical to the most fundamental:
non-reductionist approaches adapted to complex problems

unexpected action mode of sorafenib phosphatases
activate the pERK degradation

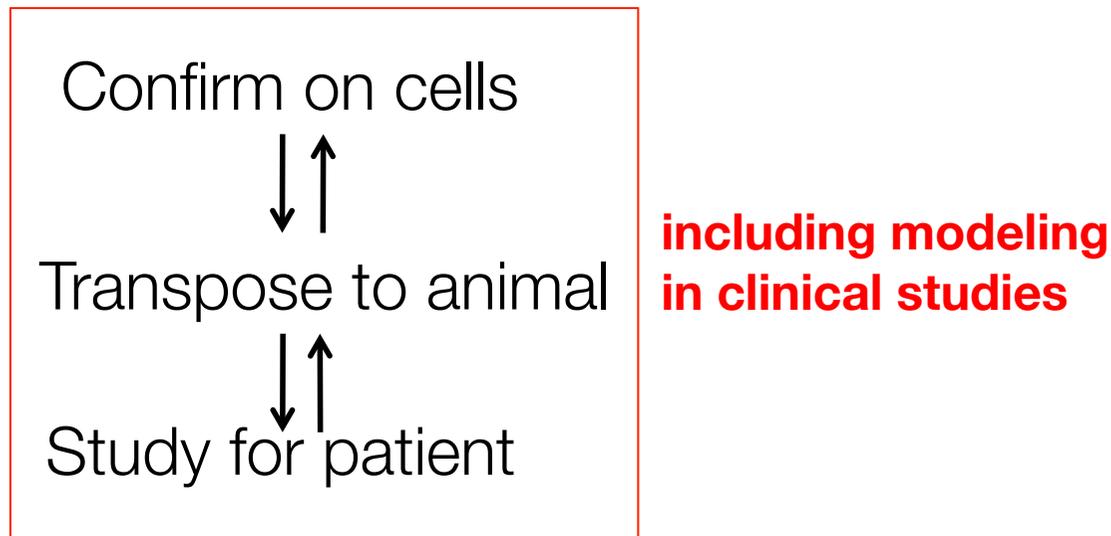
Each tumor is different → Adjust individually the treatment of the patient

Towards a predictive oncology

Most anti-cancer drugs fail in Phase 2 clinical development

Better design clinical trials to increase the chances of success and promote therapeutic innovation

Need to develop tools to integrate this complexity and to predict the impact of each therapeutic intervention



Thank for your attention

