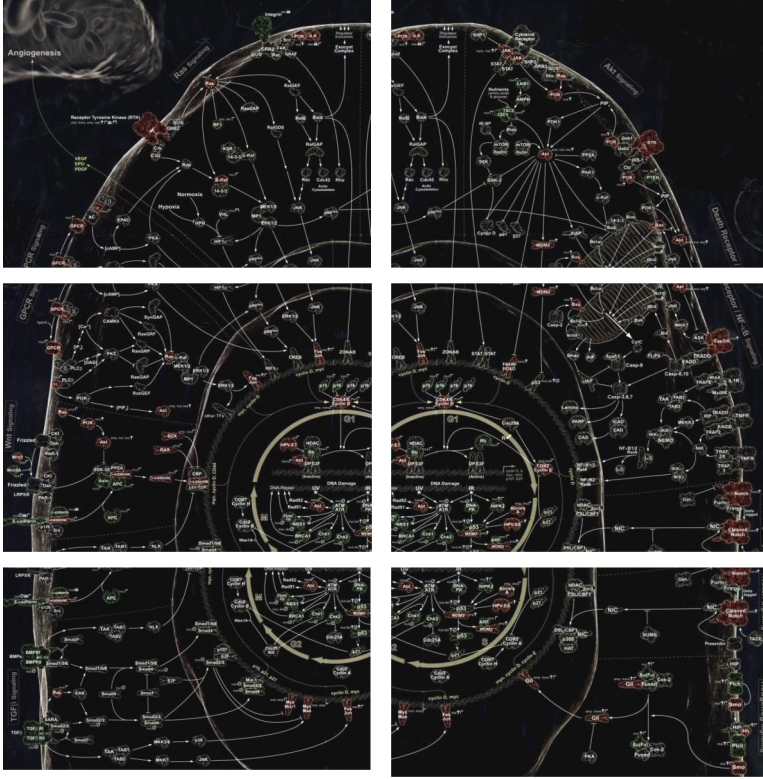




FREQUENTLY ALTERED PATHWAYS IN CANCER

CANCER BIOLOGY

AGENDA

Nov 5th: Cancer genomics- mutations

Nov 11th: Cancer genomics-copy number alteration, heterogeneity, evolution
(*recording*)

Nov 18th: Cancer Epigenetics- chromatin 3D structure, cell plasticity

Nov 25th: – Major signaling pathways leading to cancer

Dec 2th: Cancer Therapies – chemo and targeted therapies

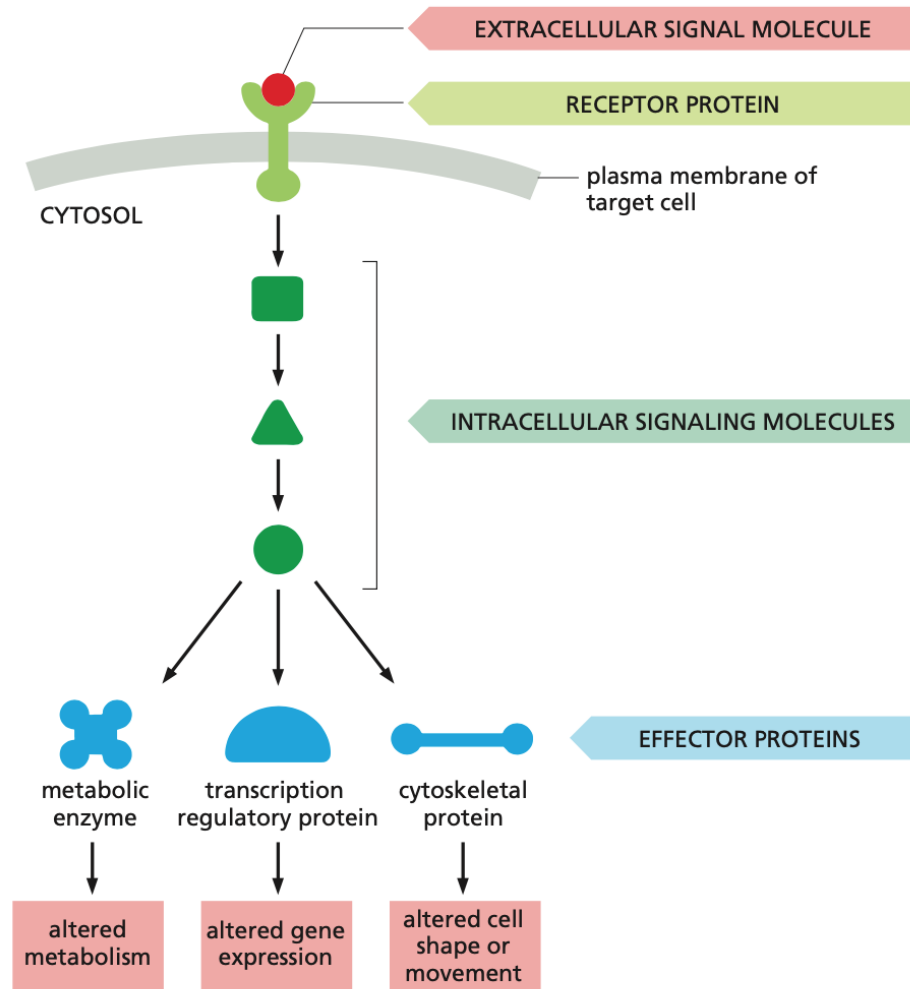
Dec 9th: Introduction to immunotherapies –

Dec 16th: questions for the exam

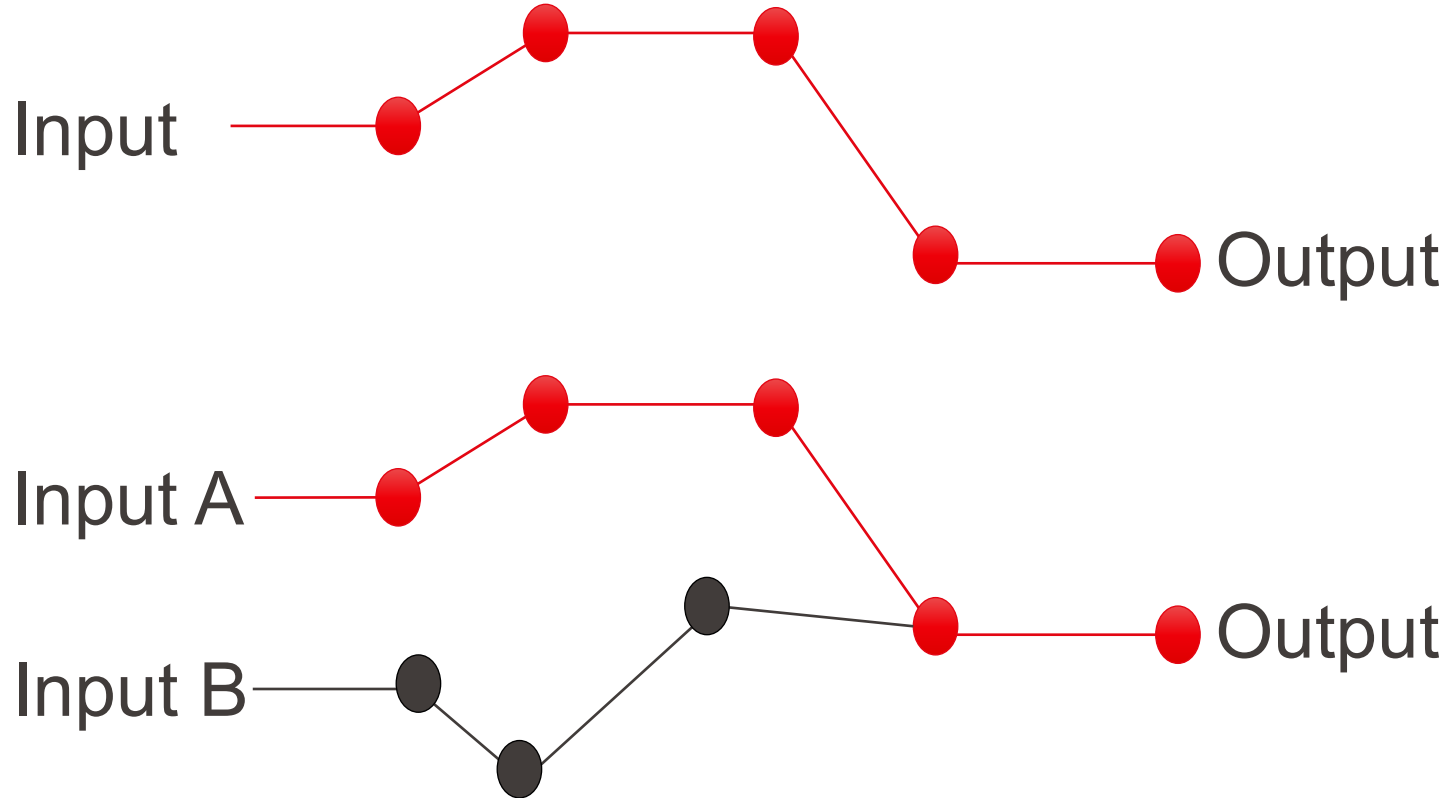
Dec 18th: Exam 1.30 PM-3.30 PM

What is a pathway

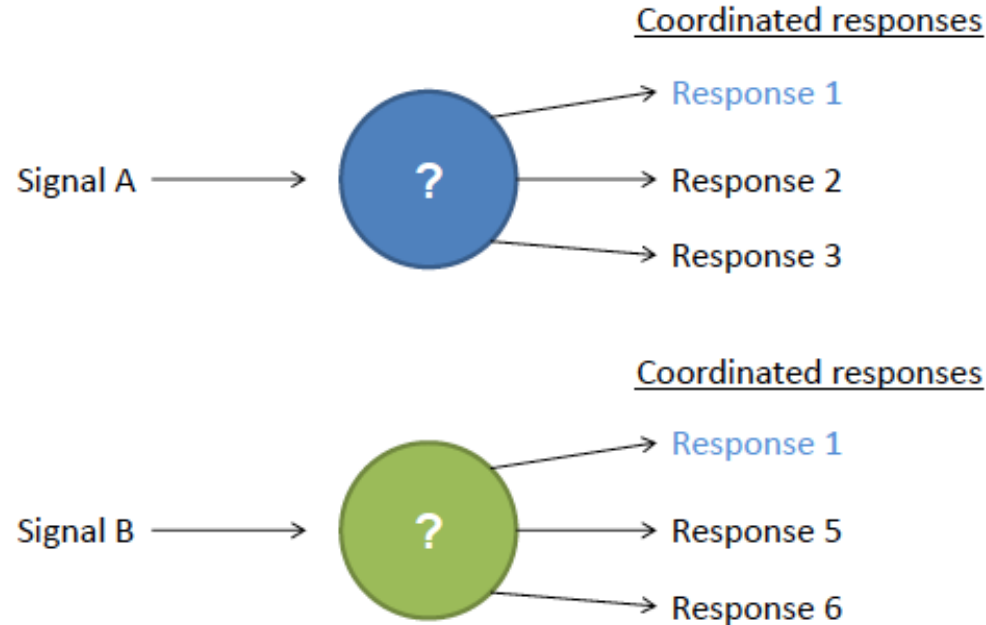
- A group of molecules that work together to coordinate a response
- The pathway works as a cascade signals
- The first molecule of the pathway receives the input and starts a cascade of events that produces output (response)



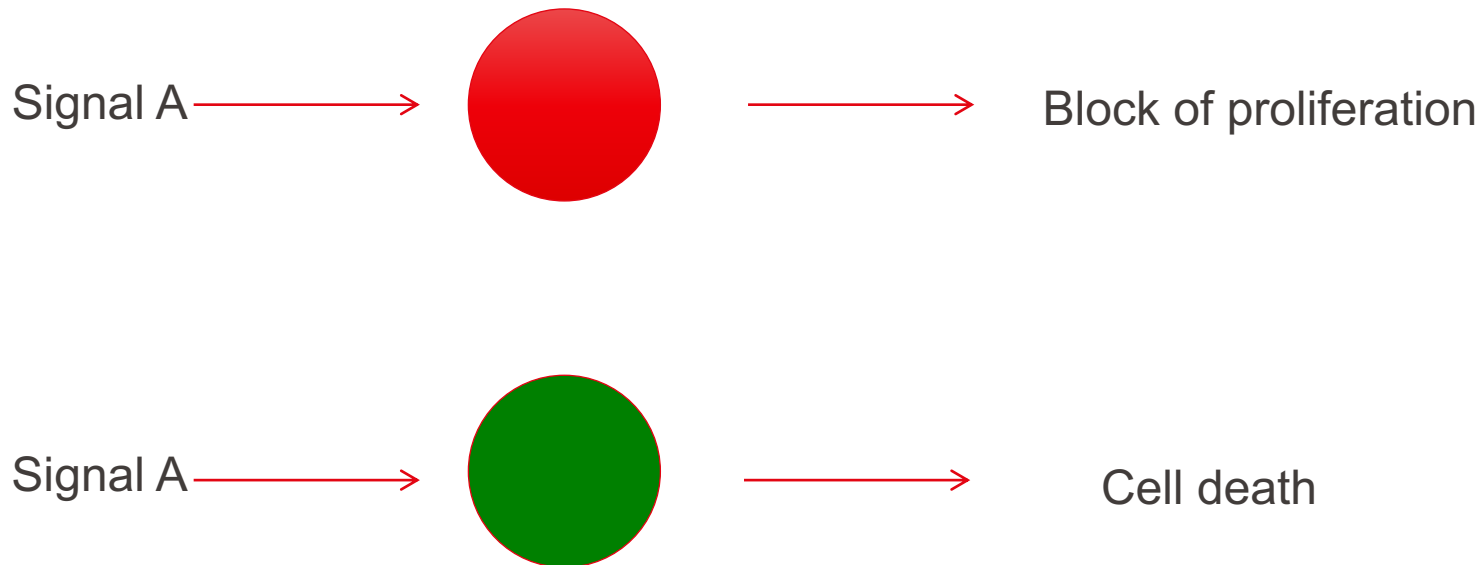
Input and Output are coordinated



Signal can coordinate multiple responses



Response can be context dependent



Difference in response can be linked to mutations in the pathway and parallel signals

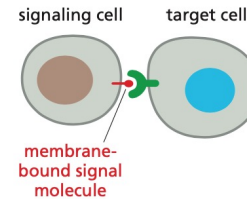
Major signaling pathways in cancer

- *Ras Signaling*
- *AKT Signaling*
- *Notch Signaling*
- *Hedgehog Signaling*
- *TGF- β Signaling*
- *WNT Signaling*
- *Cell Death Signaling*

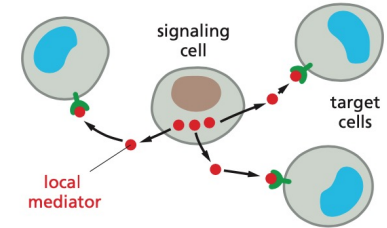
Where does the signal (=input) come from?

- *Endocrine signaling (long distances)*
- *Paracrine signaling (local mediators)*
- *Autocrine signaling*
- *Gap junction signaling (contact-dependent)*
- *Synaptic signaling (in neurons)*

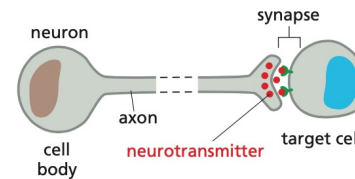
(A) CONTACT-DEPENDENT



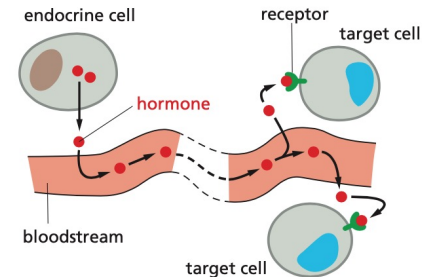
(B) PARACRINE



(C) SYNAPTIC

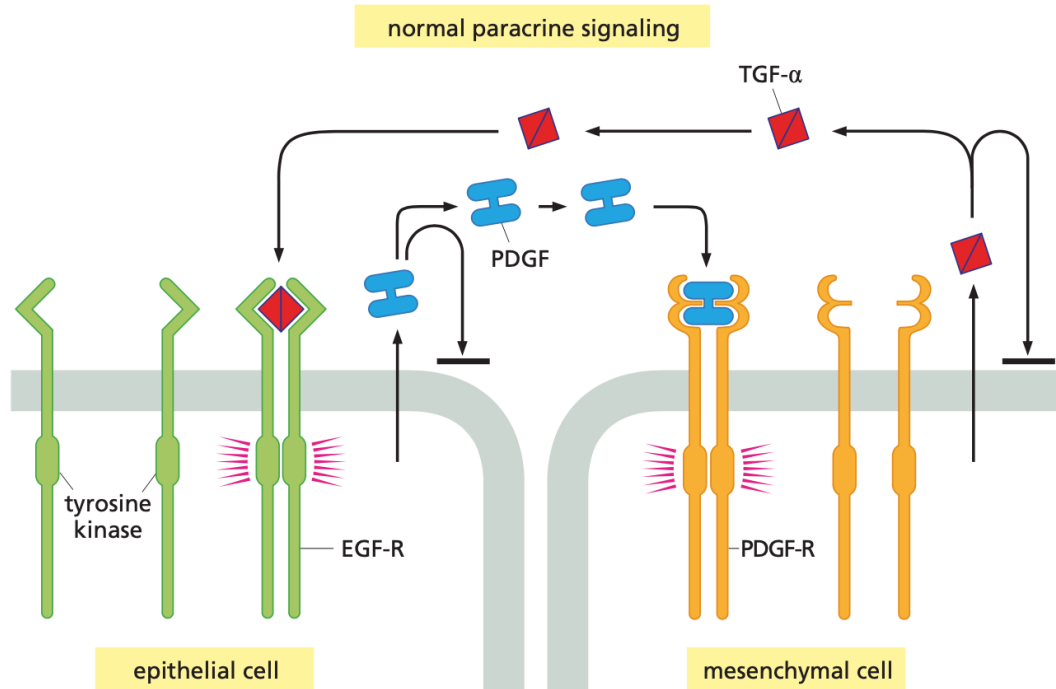


(D) ENDOCRINE



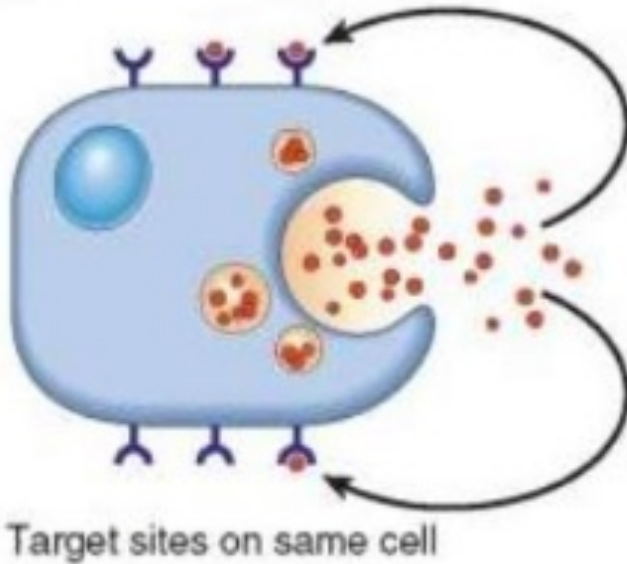
Paracrine signals

The signal molecule (e.g. interleukins, hormones) is produced by different cell types

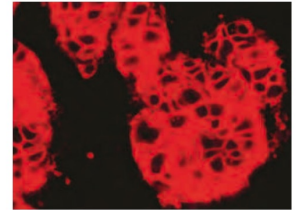


Autocrine signals

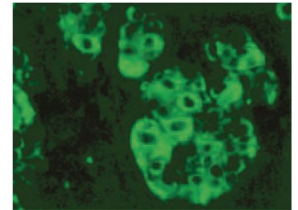
The signal molecule is produced by the same cells that received the signal



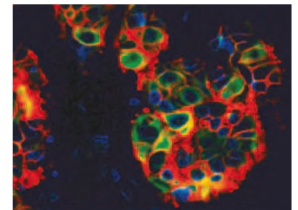
An example of autocrine signaling in successive sections of an invasive human breast carcinoma.



EGF-R



TGF- α



merged

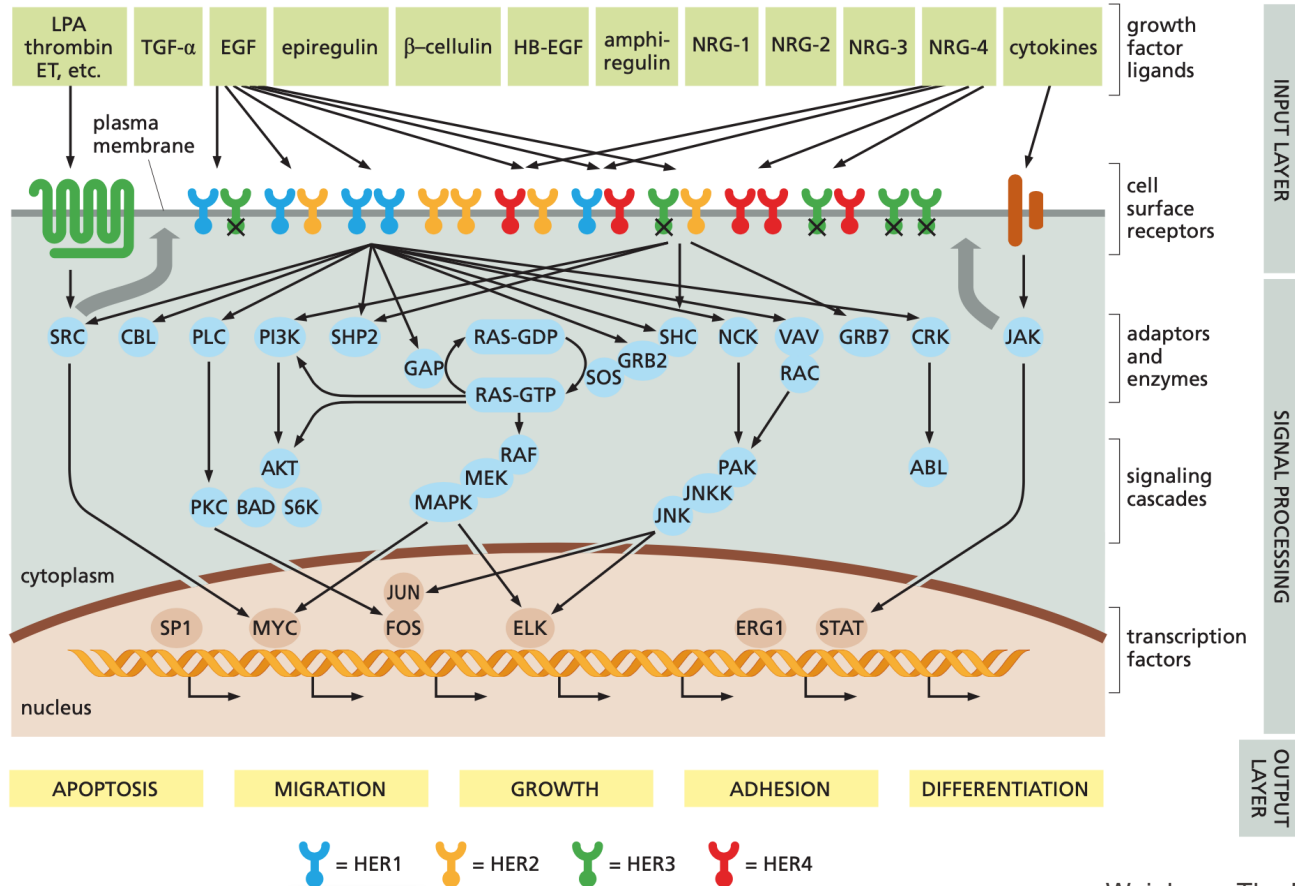
Autocrine molecules

Table 5.3 Examples of human tumors making autocrine growth factors

Ligand	Receptor	Tumor type(s)
HGF	Met	miscellaneous endocrinal tumors, invasive breast and lung cancers, osteosarcoma
IGF-2	IGF-1R	colorectal
IL-6	IL-6R	myeloma, HNSCC
IL-8	IL-8R A	bladder cancer
NRG	ErbB2 ^a /ErbB3	ovarian carcinoma
PDGF-BB	PDGF-R α/β	osteosarcoma, glioma
PDGF-C	PDGF-R α/β	Ewing's sarcoma
PRL	PRL-R	breast carcinoma
SCF	Kit	Ewing's sarcoma, SCLC
VEGF-A	VEGF-R (Flt-1)	neuroblastoma, prostate cancer, Kaposi's sarcoma
TGF- α	EGF-R	squamous cell lung, breast and prostate adenocarcinoma, pancreatic, mesothelioma
GRP	GRP-R	small-cell lung cancer

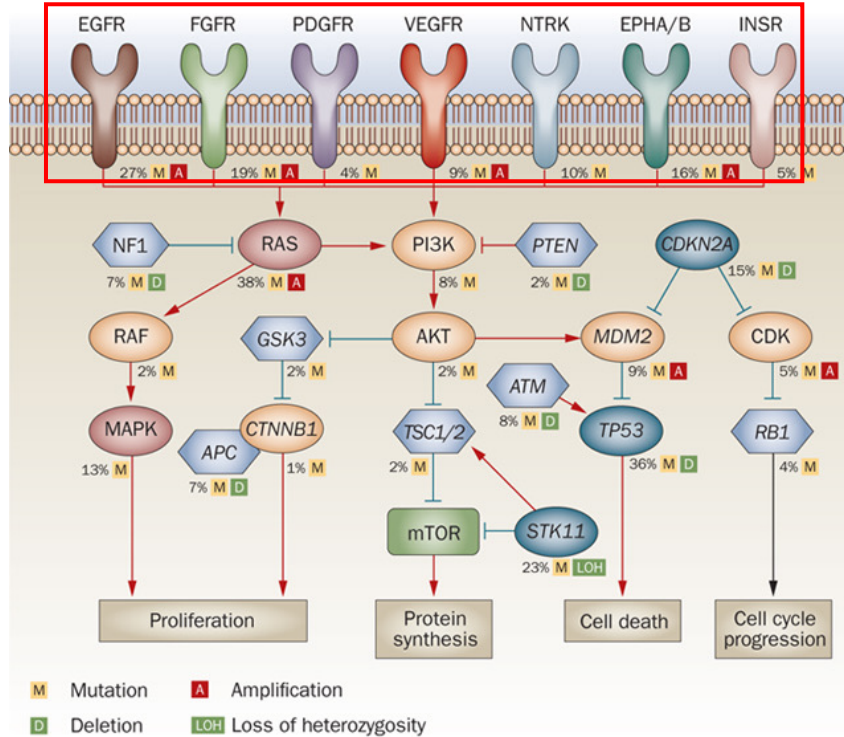
^aAlso known as HER2 or Neu receptor

One ligand can bind to multiple receptors



Receptor Tyrosine Kinases

Cell growth



Epidermal growth factor

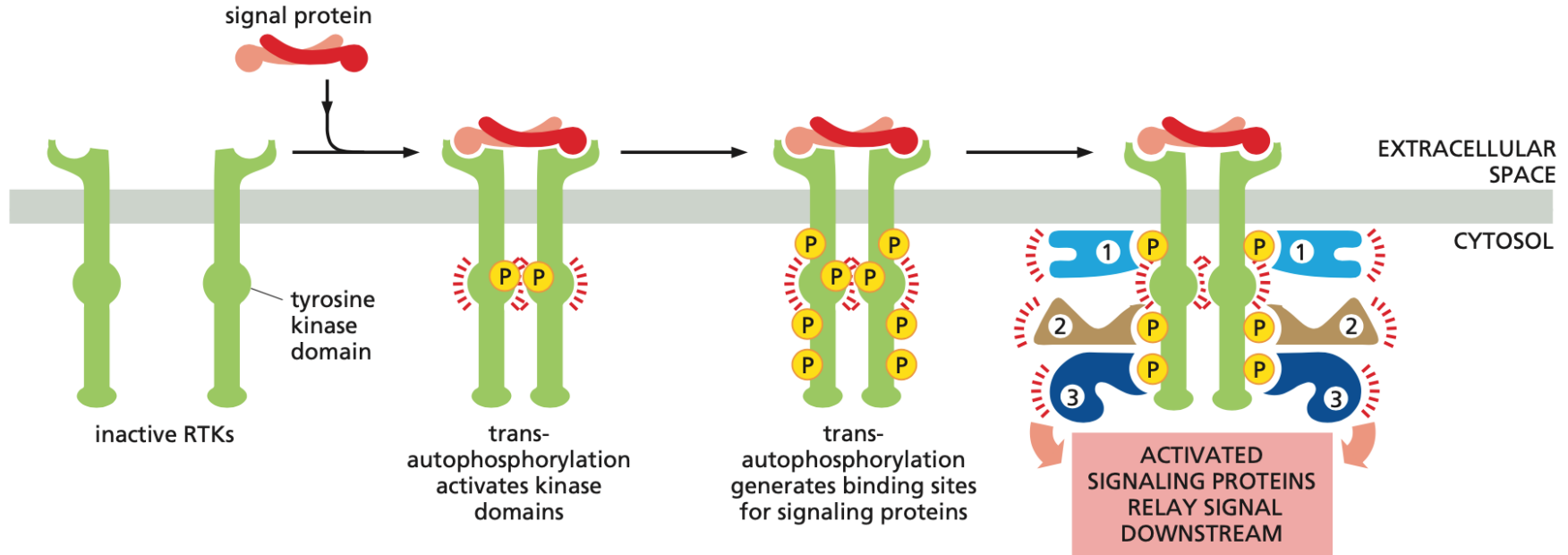
Fibroblast growth factor

Platelet-derived growth factor

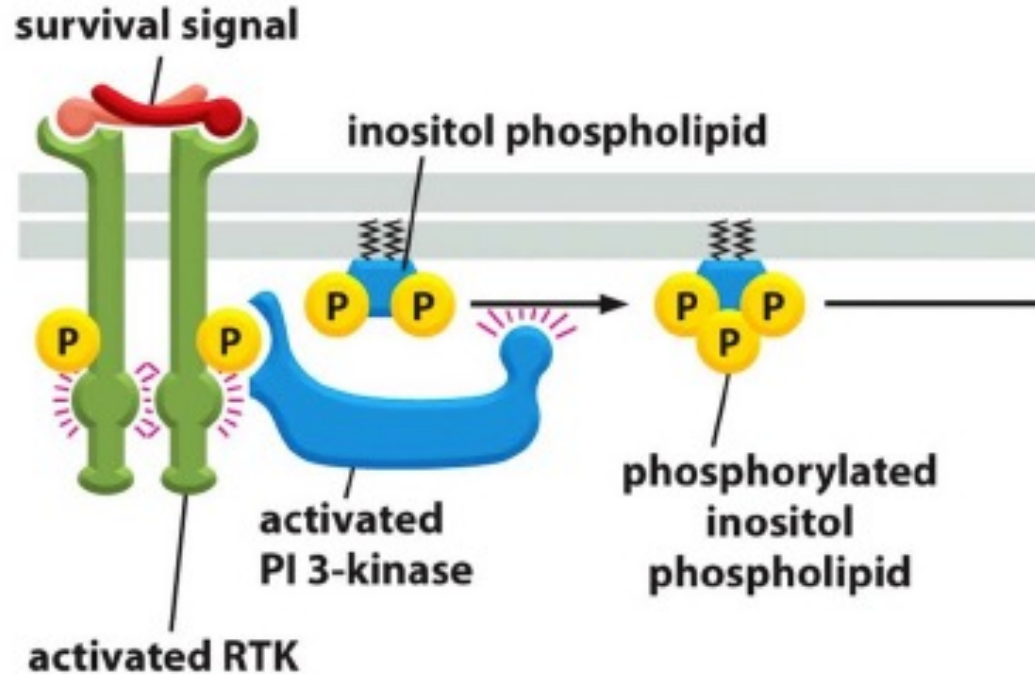
Vascular endothelial growth factor

Insulin growth factor

Receptor Tyrosine Kinases dimerization

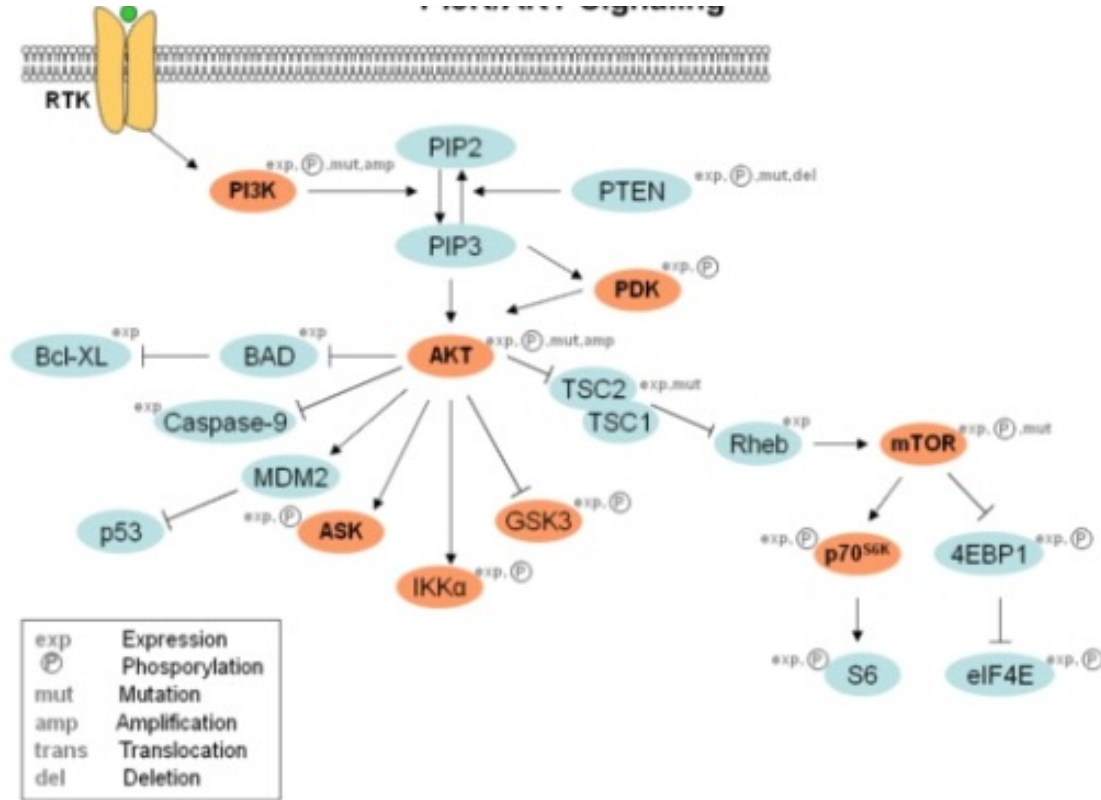


Downstream RTK proteins: the case of PI3K

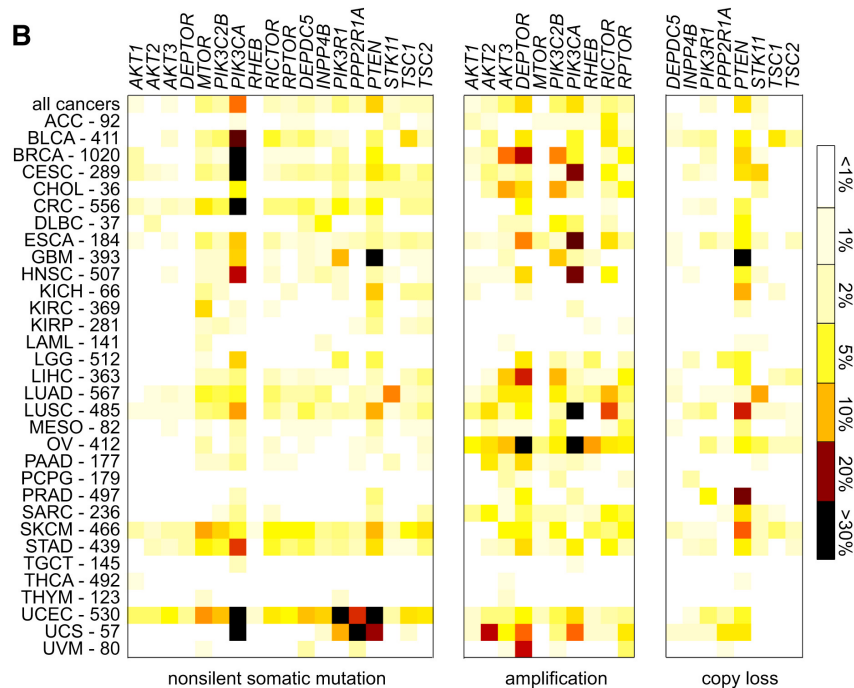
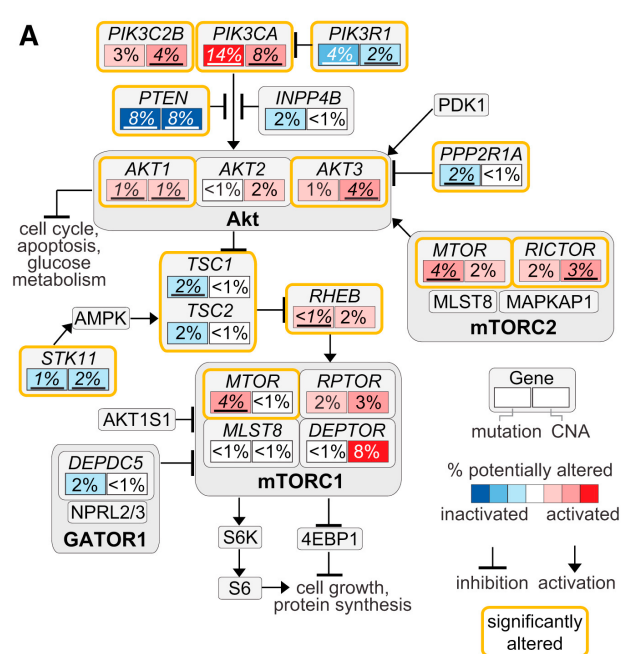




PI3-kinase activates Akt

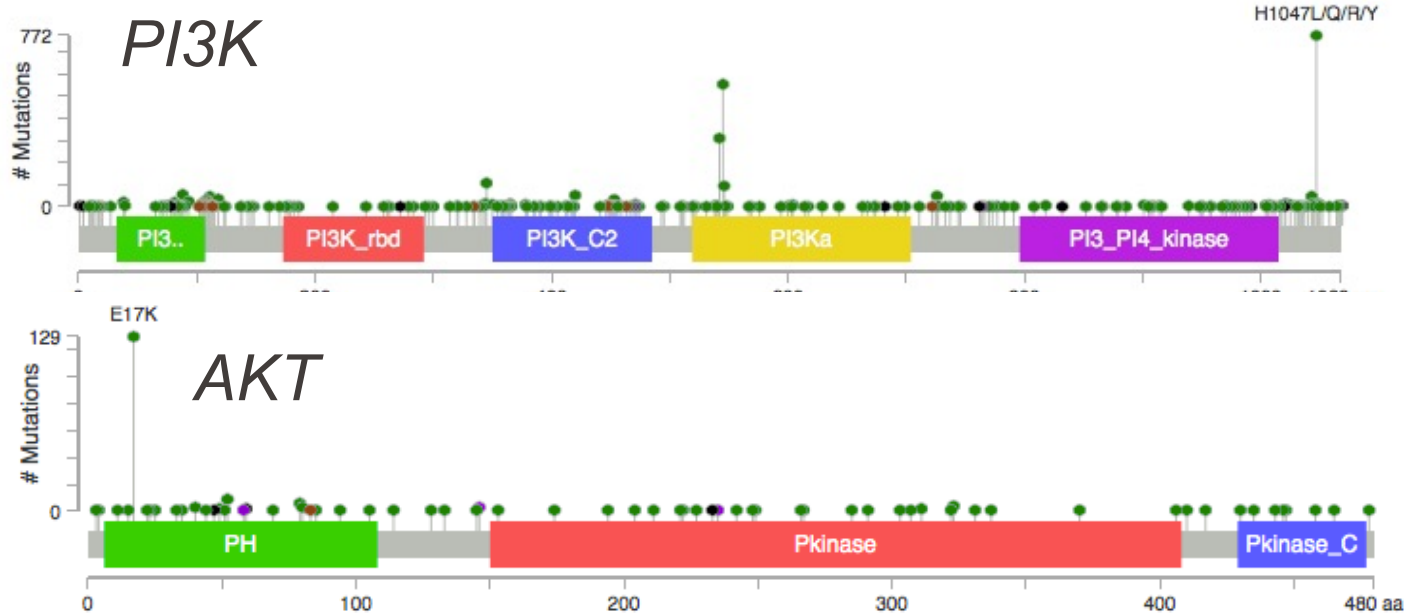


PI3K/Akt pathway is frequently altered across tumors

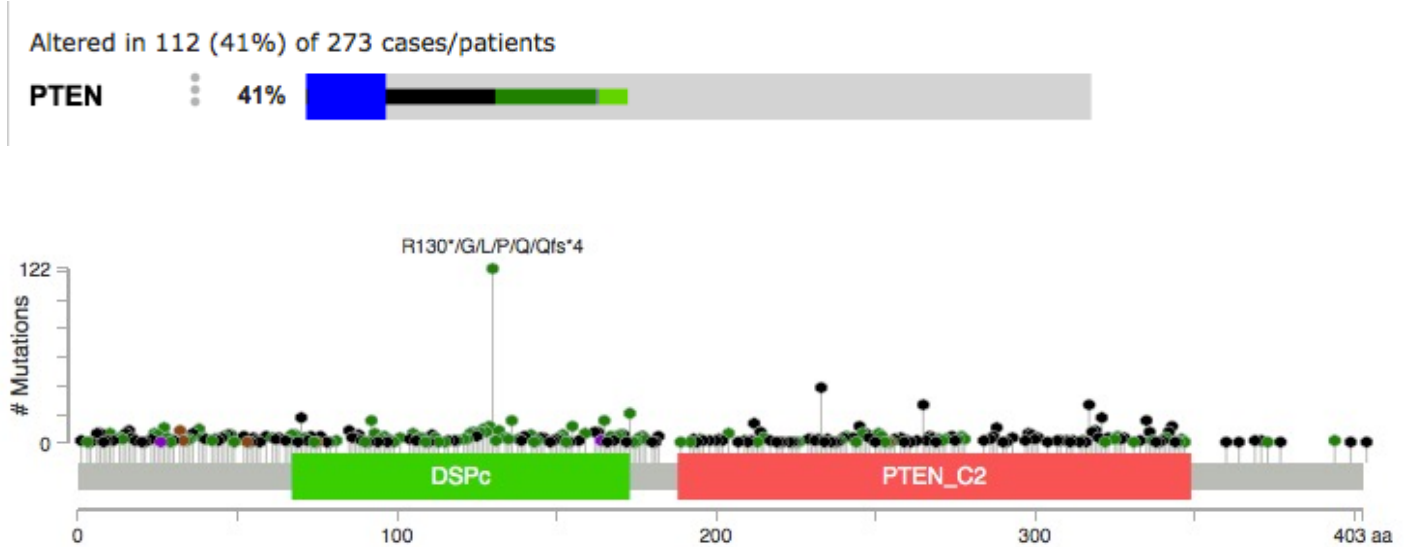


LAML, Acute Myeloid Leukemia; ACC, Adrenocortical carcinoma; BLCA, Bladder Urothelial Carcinoma; LGG, Brain Lower Grade Glioma; BRCA, Breast invasive carcinoma; CESC, Cervical squamous cell carcinoma and endocervical adenocarcinoma; CHOL, Cholangiocarcinoma; CRC, Colorectal adenocarcinoma (combining COAD and READ projects); ESCA, Esophageal carcinoma; GBM, Glioblastoma multiforme; HNSC, Head and Neck squamous cell carcinoma; KICH, Kidney Chromophobe; KIRC, Kidney renal clear cell carcinoma; KIRP, Kidney renal papillary cell carcinoma; LIHC, Liver hepatocellular carcinoma; LUAD, Lung adenocarcinoma; LUSC, Lung squamous cell carcinoma; DLBC, Lymphoid Neoplasm Diffuse Large B-cell Lymphoma; MESO, Mesothelioma; OV, Ovarian serous cystadenocarcinoma; PAAD, Pancreatic adenocarcinoma; PCPG, Pheochromocytoma and Paraganglioma; PRAD, Prostate adenocarcinoma; SARC, Sarcoma; SKCM, Skin Cutaneous Melanoma; STAD, Stomach adenocarcinoma; TGCT, Testicular Germ Cell Tumors; THYM, Thymoma; THCA, Thyroid carcinoma; UCS, Uterine Carcinosarcoma; UCEC, Uterine Corpus Endometrial Carcinoma.

Hot-spot mutations

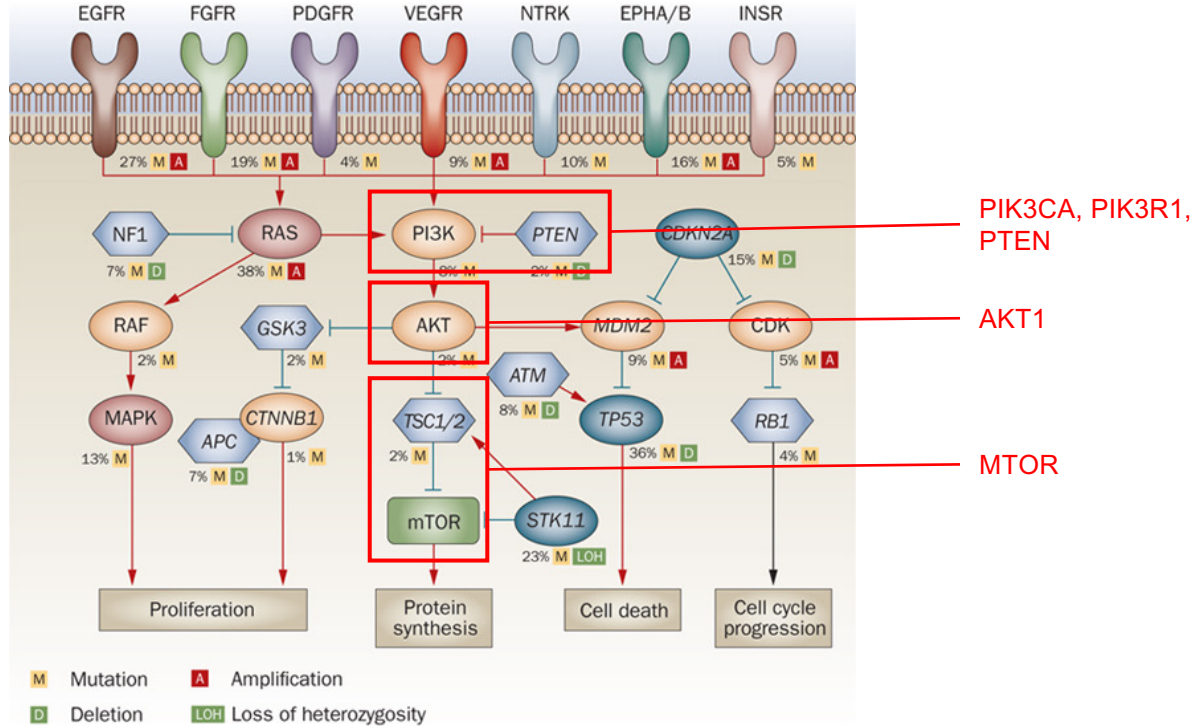


PTEN is deleted or mutated in several cancers



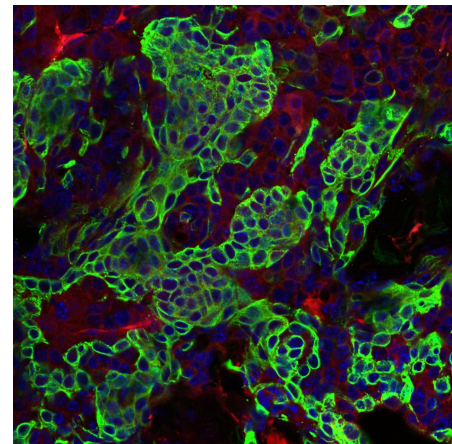
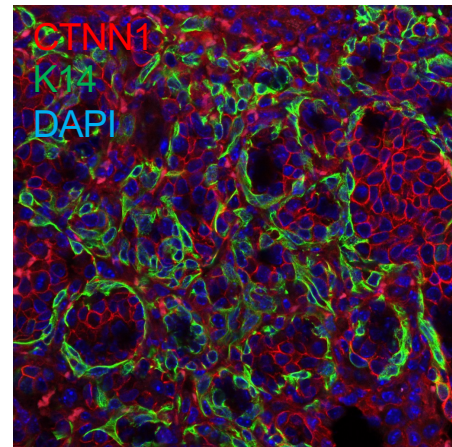
PI3K/Akt pathway

■ Survival & Translation

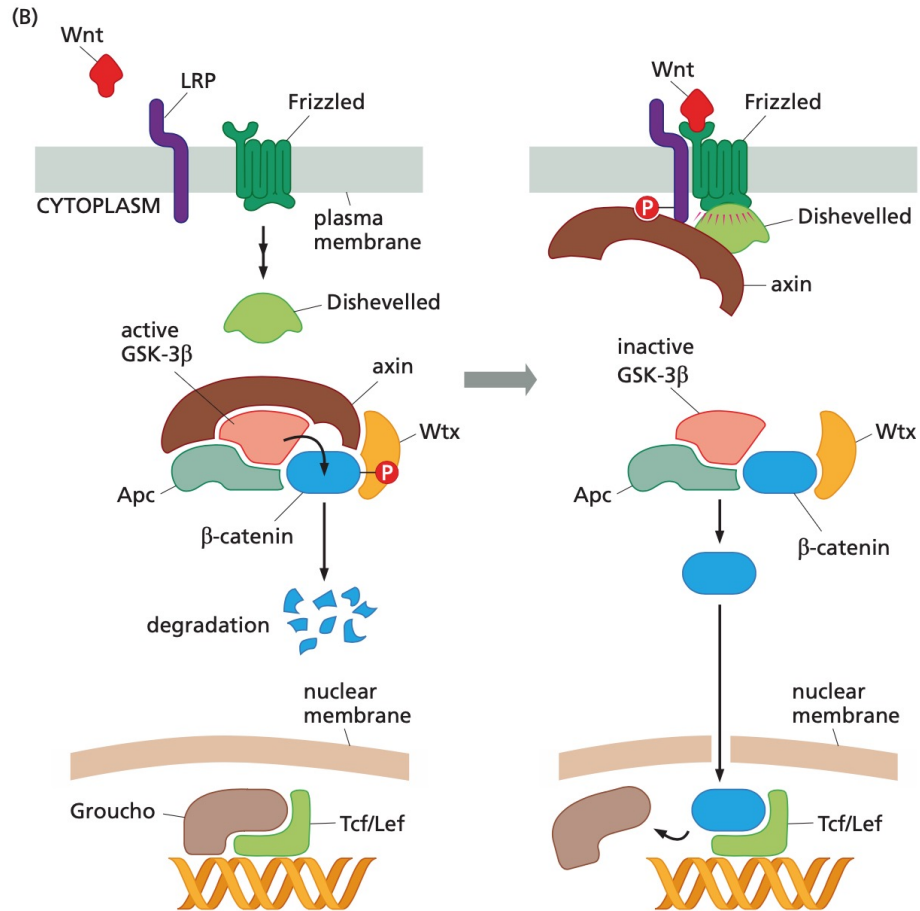
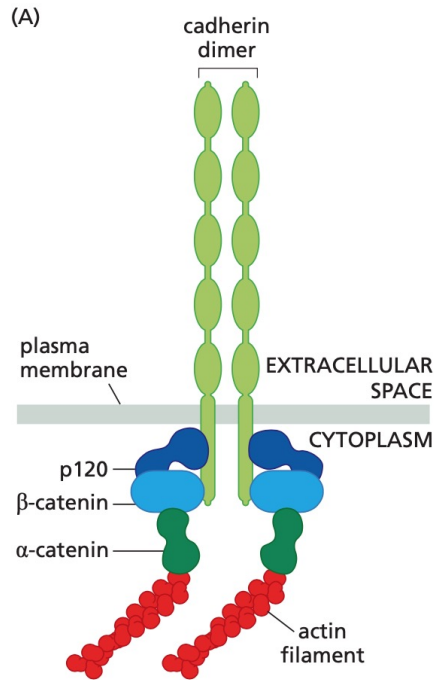


The Wnt pathway

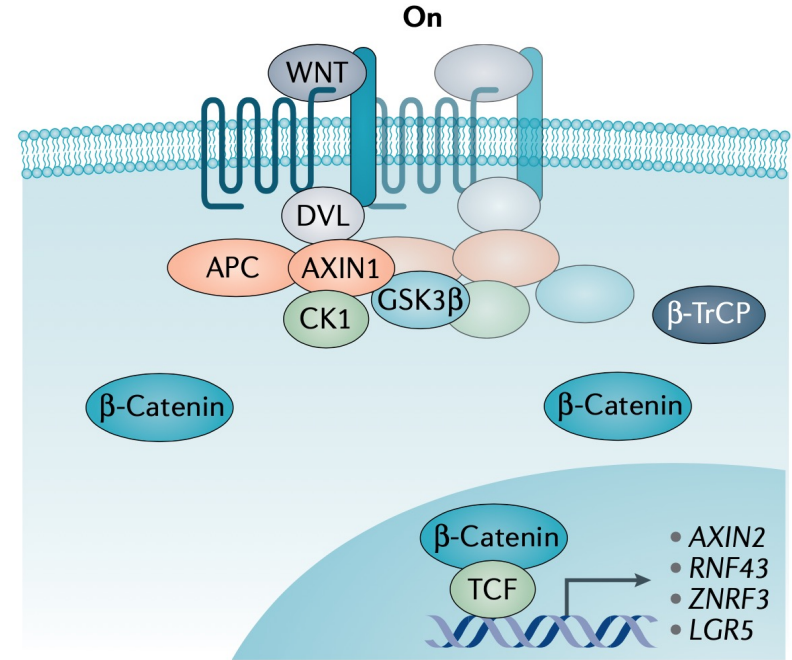
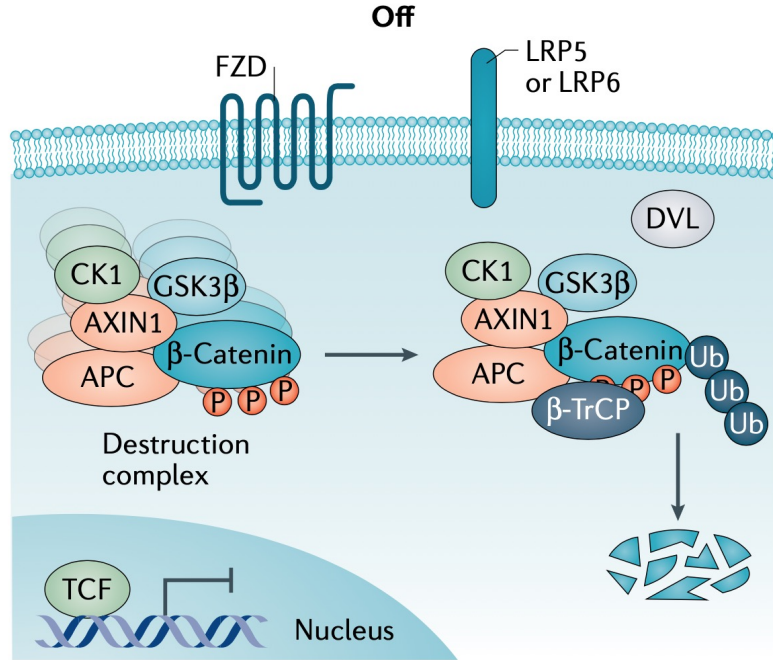
- Important during development (patterning decisions).
- 19 ligands (Wnt members)
- More than 15 receptors: Frizzled, LRP5, LRP6, RYK,...
- Canonical and non-canonical signaling.
- Roles in cancer:
 - Cancer stem cells: e.g. self-renewal, TERT expression.
 - Invasion and metastasis through EMT.



The Wnt pathway



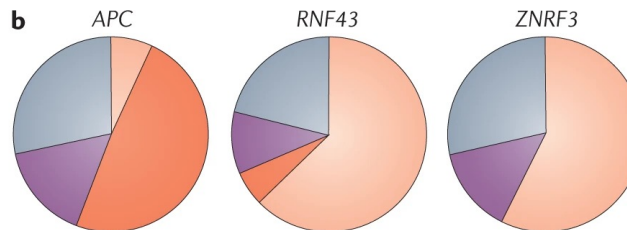
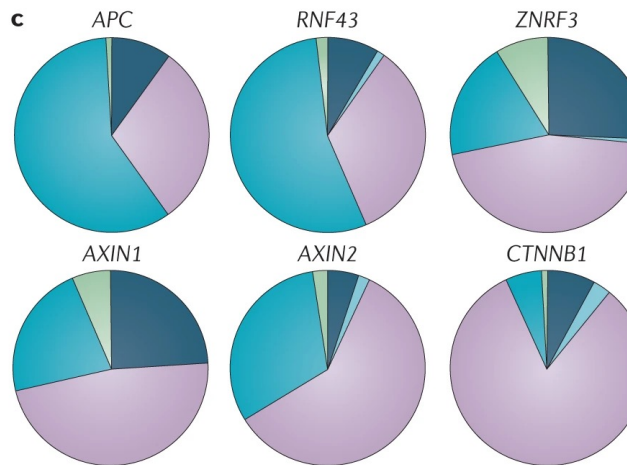
Canonical Wnt signaling



Alterations of the Wnt pathway in cancer

a

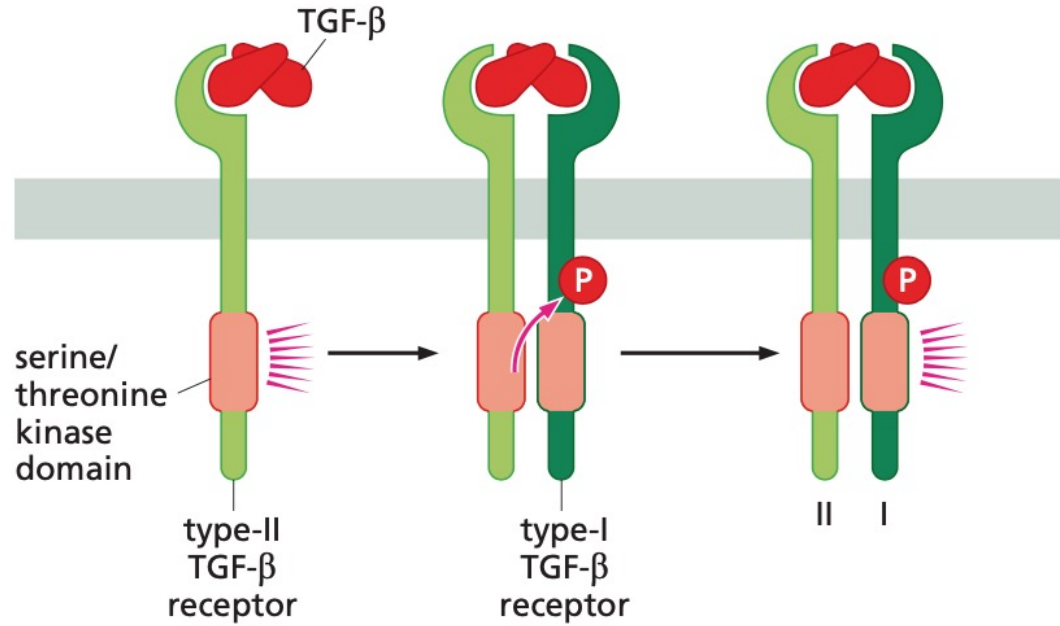
APC	Colorectal adenocarcinoma	66.55%
	Uterine corpus endometrial carcinoma	14.34%
	Skin cutaneous melanoma	14.25%
	Stomach adenocarcinoma	13.38%
	Bladder urothelial carcinoma	7.04%
RNF43	Uterine corpus endometrial carcinoma	15.28%
	Stomach adenocarcinoma	10.20%
	Colorectal adenocarcinoma	7.90%
	Pancreatic ductal adenocarcinoma	5.95%
ZNRF3	Adenocortical carcinoma	20.43%
	Uterine corpus endometrial carcinoma	6.23%
	Skin cutaneous melanoma	5.35%
AXIN1	Liver hepatocellular carcinoma	7.77%
	Uterine corpus endometrial carcinoma	6.42%
AXIN2	Uterine corpus endometrial carcinoma	7.36%
	Colorectal adenocarcinoma	5.04%
CTNNB1	Liver hepatocellular carcinoma	25.47%
	Uterine corpus endometrial carcinoma	25.47%
	Adenocortical carcinoma	15.05%
	Skin cutaneous melanoma	6.68%
	Stomach adenocarcinoma	6.68%
	Colorectal adenocarcinoma	5.71%

b

c


The TGF β pathway

- Controls: cell cycle, metastasis, invasion, angiogenesis, wound healing, cancer stem cells,...
- Cell- and context-dependent.
- Integrated in many other signaling pathways (Wnt, Notch, Ras, etc).
- Pleiotropic effects: can exert cytostatic and tumor-suppressive effects in early stage tumors, but can also induce proliferation, invasion, and angiogenesis in advanced tumors.

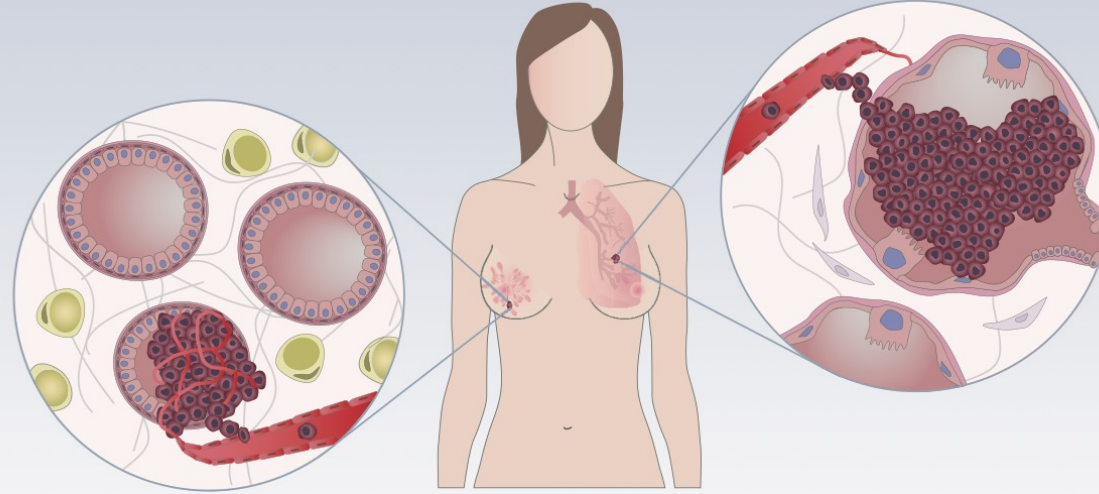
The TGF β pathway





A case of pleiotropism: The TGF β pathway

TGF- β in tumor progression and metastasis



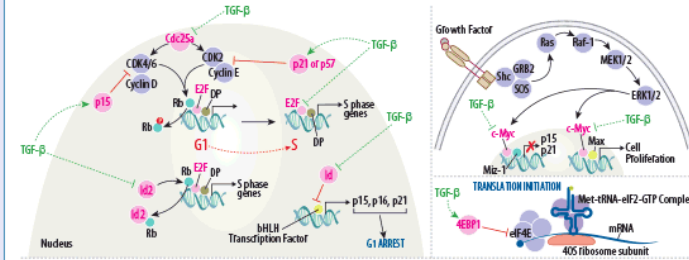
Premalignant	Malignancy	Dissemination	Metastasis
Tumor-suppressor signals: Cytostasis Differentiation Apoptosis Inflammatory suppressor Stroma response suppressor	Evasion of immunity Evasion of cytostasis Autocrine mitogenic signals Motility and migration	EMT Stromal recruitment: Fibroblasts BMD cells Metastasis endowing properties Niche preparation	Extravasation BMD cells mobilization Stroma-modifying factors

TGF β target genes

Tumor-Suppressing Effects of TGF- β

Inhibits Cell Proliferation	Induces Apoptosis	Activates Autophagy	Inhibits Growth Factors in the Tumor Stroma	Inhibits Angiogenesis	Suppresses Inflammation
TGF- β Target Genes p15, p21, p57, 4EBP1	TGF- β Target Genes BAX, BIM, DAPK, Fas, GADD45B	TGF- β Target Genes ATG5, ATG7, Beclin 1/ATG6, DAPK		TGF- β Target Genes Thrombospondin	TGF- β Target Genes FoxP3
TGF- β Target Genes CDK2A, E2F-1, M1-3, c-Myc	TGF- β Target Genes Bcl-xL, Bcl-2		TGF- β Target Genes HGF, MSP, TGF- α		TGF- β Target Genes GATA-3, T-bet

Inhibits Cell Proliferation

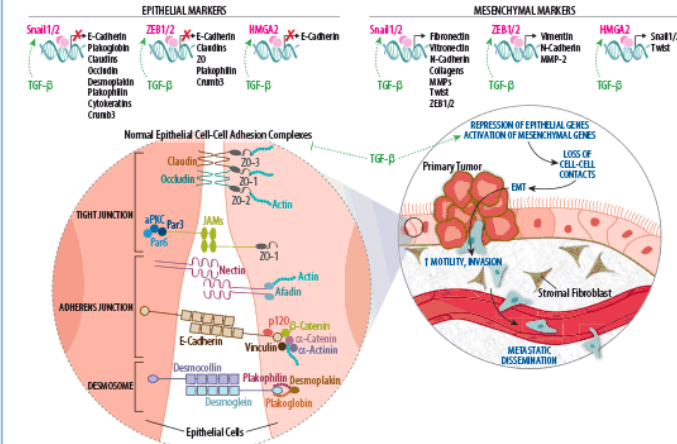


TGF- β inhibits cell proliferation by inducing the expression of 4EBP1 and the cyclin-dependent kinase (CDK) inhibitors, p15, p21, and p57. 4EBP1 binds to eukaryotic initiation factor 4E (eIF4E) and inhibits protein translation (bottom right panel), while p15, p21, and p57 prevent cell cycle progression by inhibiting the activities of CDK-cyclin complexes that are required for the G1/S transition (left panel). Additionally, TGF- β represses the expression of CDK2A phosphatase, which is also required for CDK-cyclin activation (left panel), and negatively regulates the expression of multiple other factors involved in driving cell cycle progression and cell proliferation, including the Id proteins (left panel), E2F (left panel), and c-Myc (top right panel).

Tumor-Promoting Effects of TGF- β

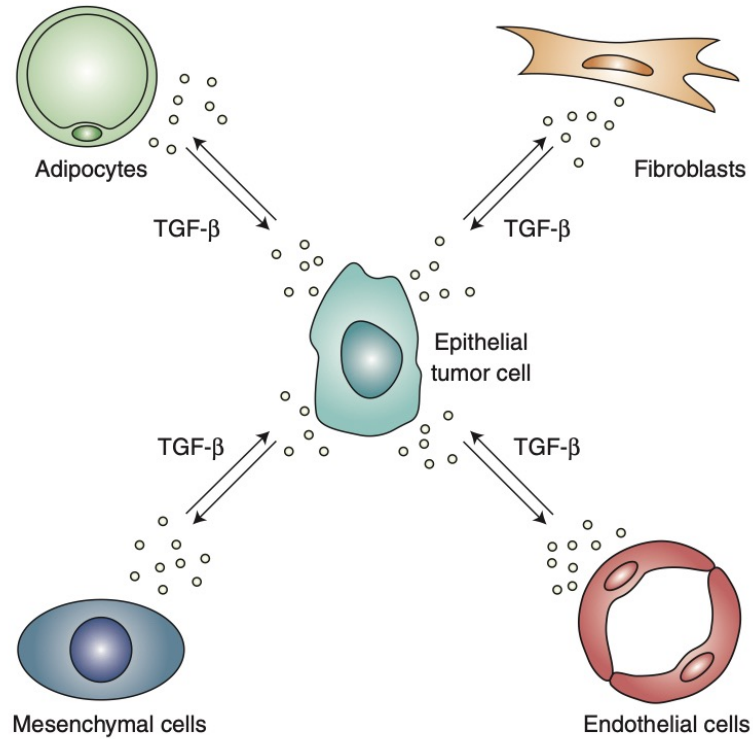
Promotes Cell Proliferation	Suppresses the Immune Response	Promotes Angiogenesis	Promotes Cancer Stem Cell Self-Renewal	Promotes the Epithelial-to-Mesenchymal Transition	Promotes Metastasis
TGF- β Target Genes PDGF- β	TGF- β Target Genes FoxP3	TGF- β Target Genes VEGF, MMP-2, MMP-9	TGF- β Target Genes LIF, SOX4	TGF- β Target Genes Snail1/2, ZEB1/2, HMG2	TGF- β Target Genes HDM2, MMP-9
	TGF- β Target Genes Fas Ligand, GATA-3, Granzyme A/B, IFN- γ , MICA, HMG20, Nrp30, Perforin, T-bet				

Promotes the Epithelial-to-Mesenchymal Transition



TGF- β signaling in epithelial tumor cells promotes an epithelial-to-mesenchymal transition by inducing the expression of transcription factors, such as Snail1/2, ZEB1/2, and HMG2, which repress the expression of epithelial cell adhesion proteins and induce the expression of mesenchymal proteins. These changes promote the loss of cell polarity and cell-cell contacts and the acquisition of a migratory, invasive phenotype that may allow cancer cell dissemination.

The TGF β pathway



The TGF β pathway

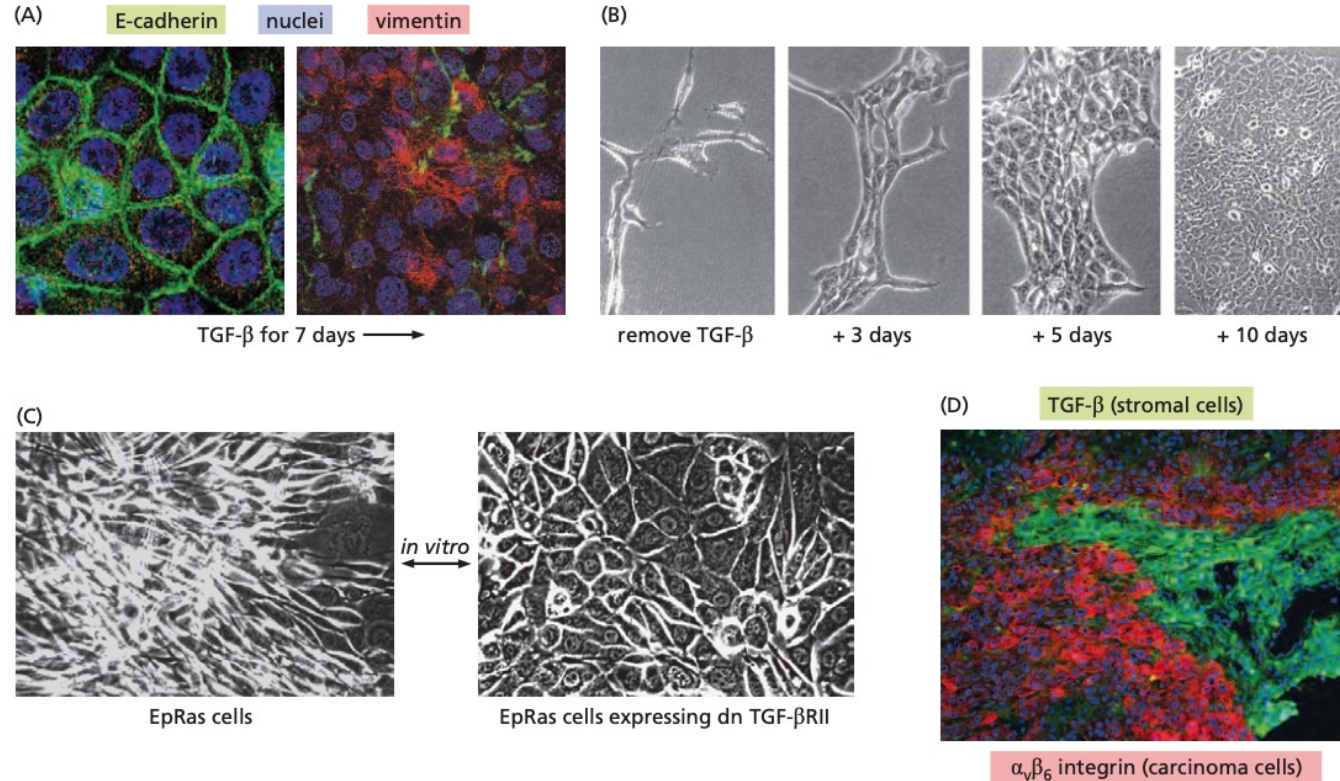
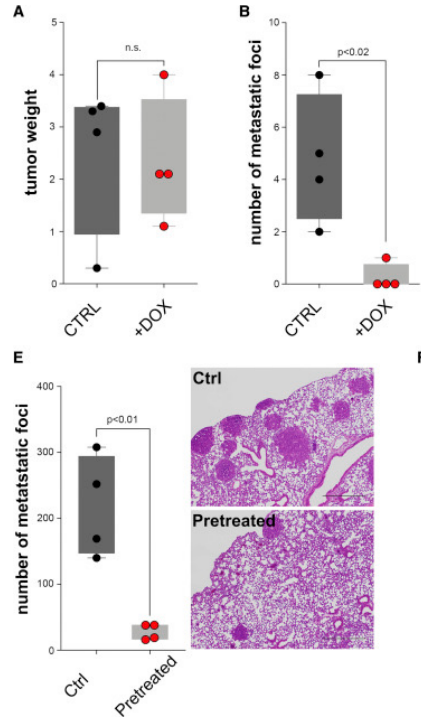


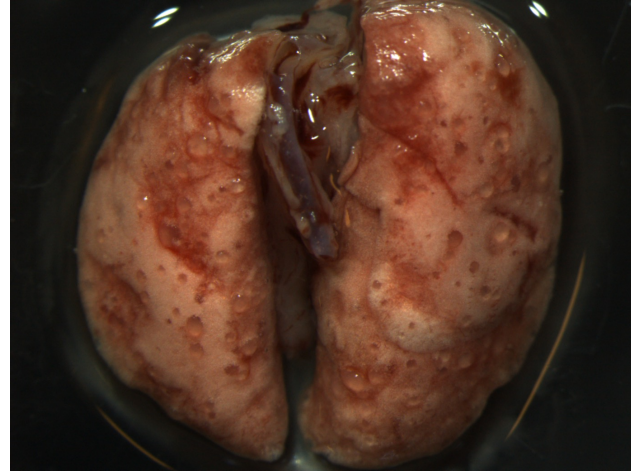
Figure 14.19 Control of the EMT by TGF- β and its effects on tumorigenic cells Immortalized mouse mammary epithelial cells

epithelial transition (MET). (C) Use of a dominant-negative (dn) type II TGF- β receptor (which effectively blocks autocrine TGF- β

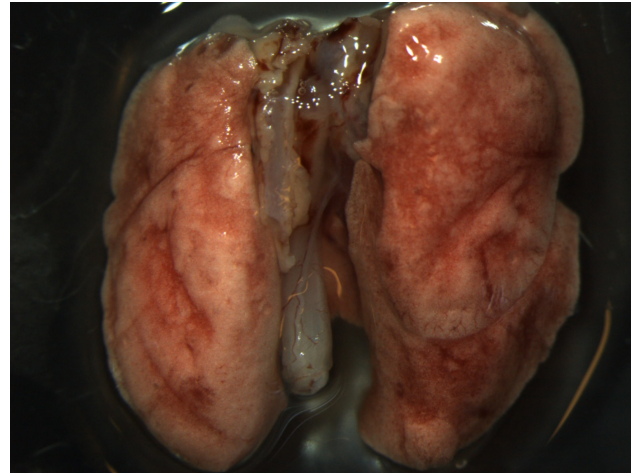
The TGF β pathway



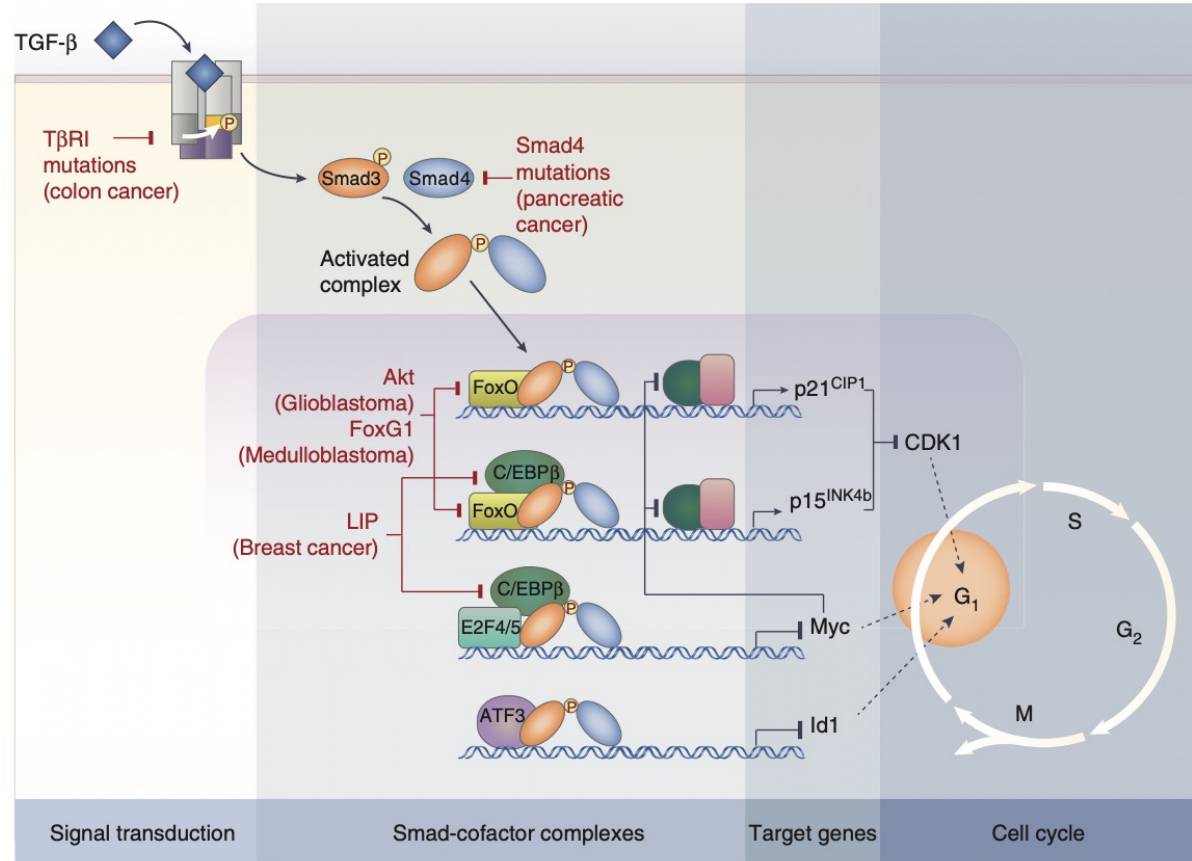
Vehicle pretreated



TGFBRI pretreated



The TGF β pathway: mutations in cancer



Metabolic alterations

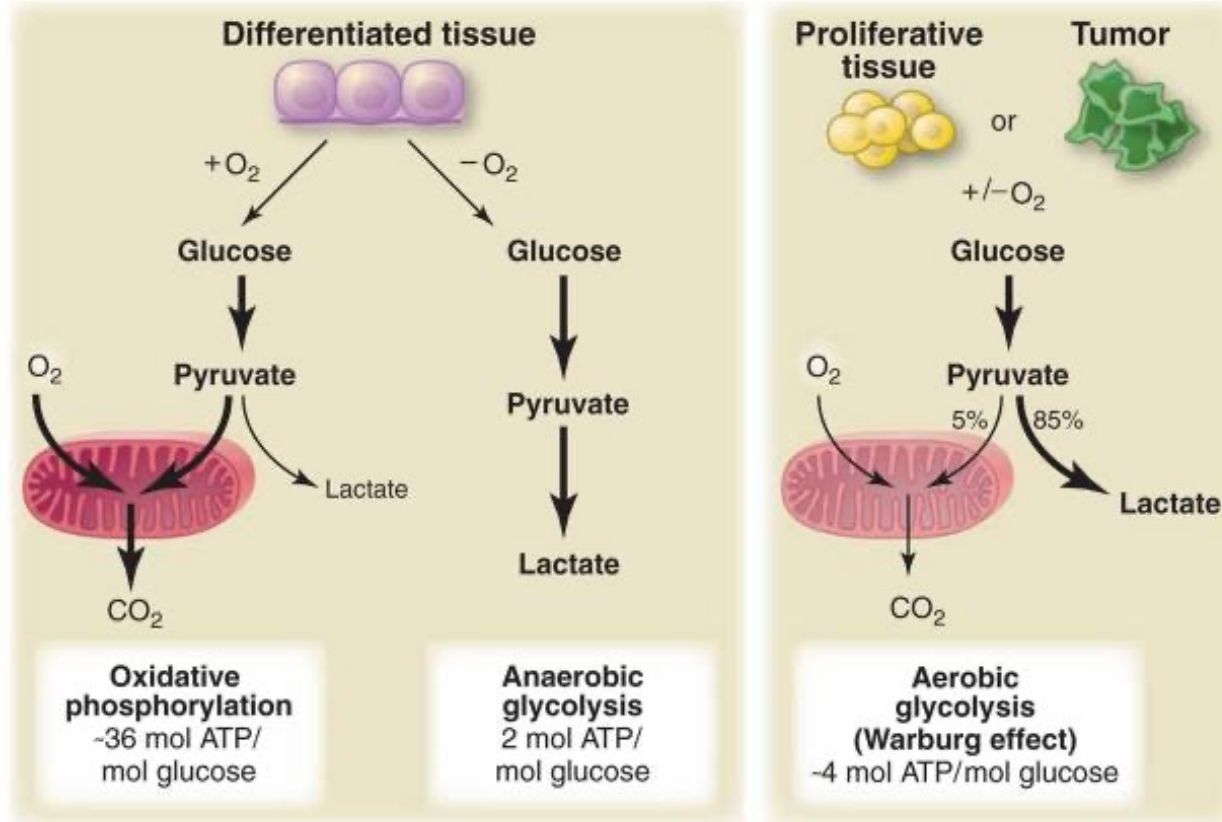
- **Change the use of metabolites (associated with cancer cell proliferation)**
- **Synthesis of new metabolites (associated with mutation in some genes)**

Metabolic alterations

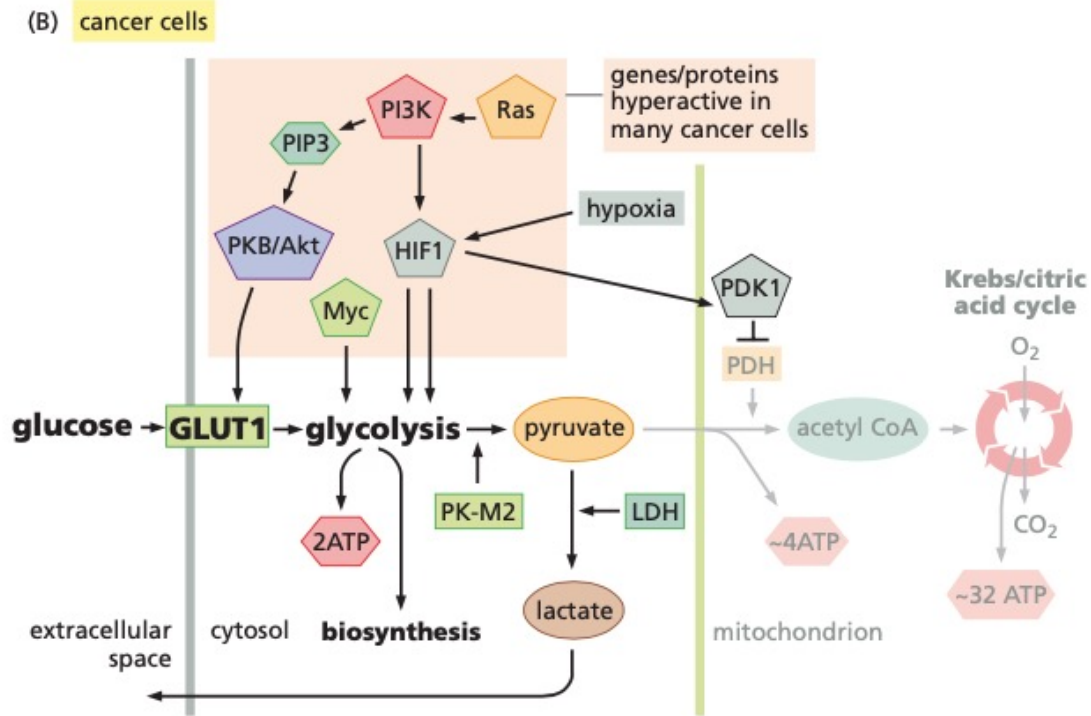
Cancer cells have a different diet compared to the normal cells



The Warburg effect

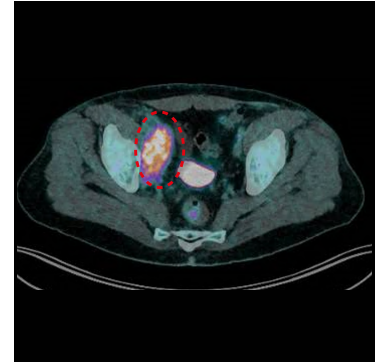


The Warburg effect

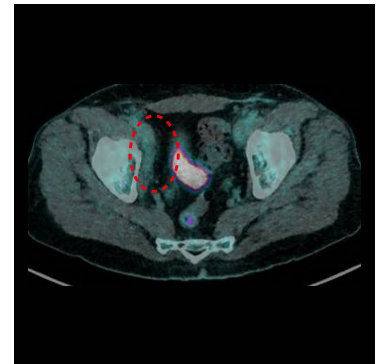


PET-CT

detection



after treatment



Production of new metabolites



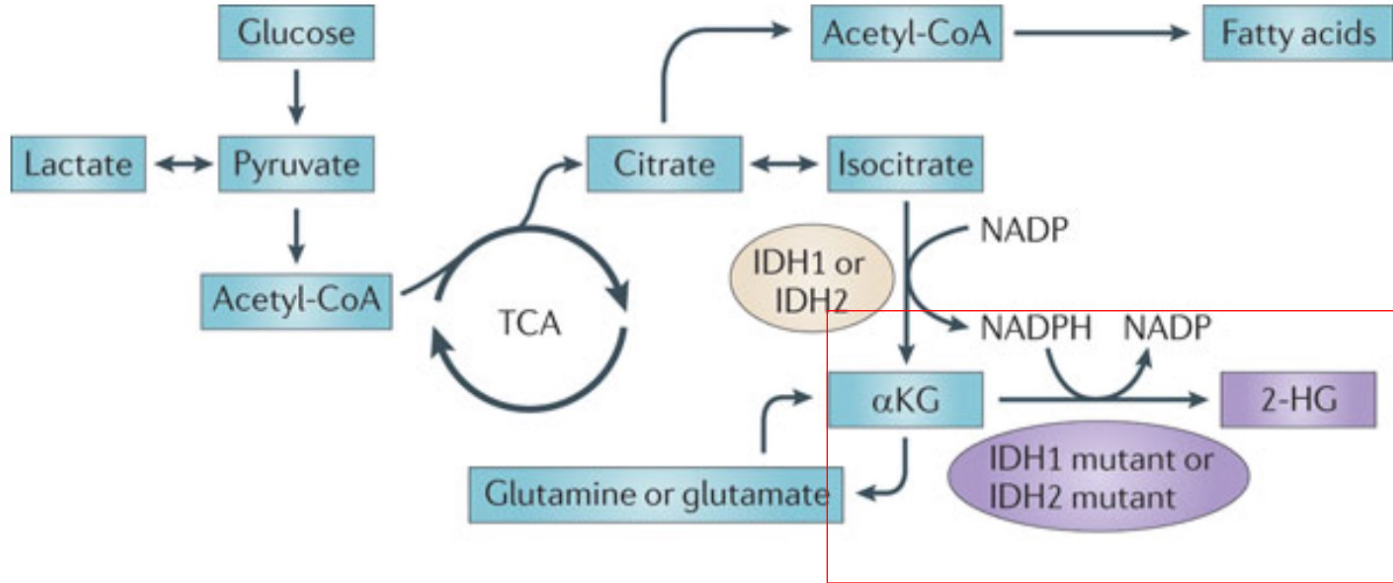
Associated with gene mutations

Mutation of IDH1 and IDH2 genes

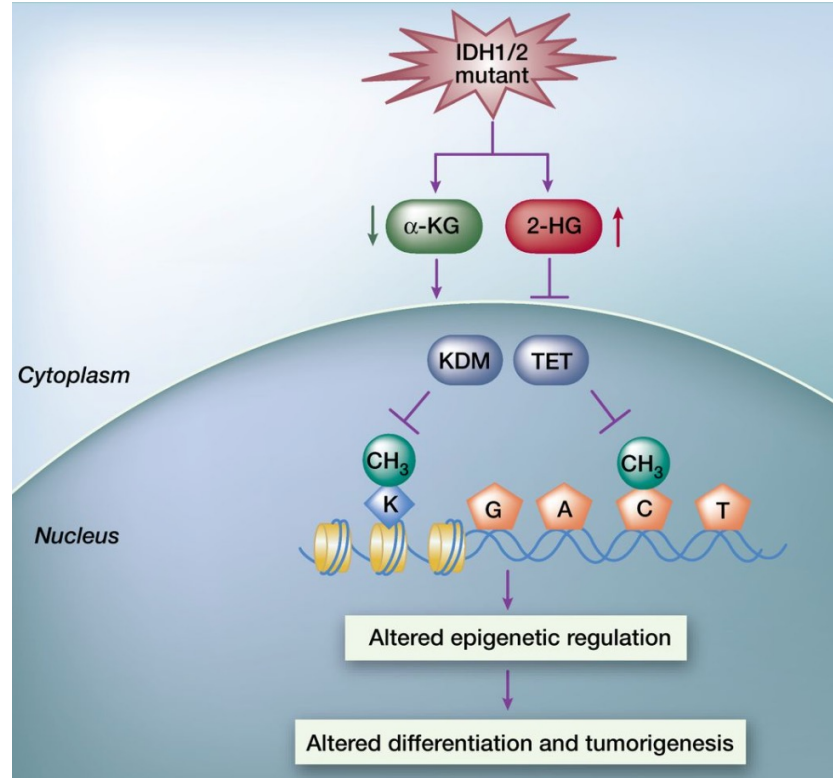
*IDH1 **R**132X (predominant in glioma; arginine R at position 132 is replaced by another amino acid, giving a missense mutation)*

*IDH2 **R**140X (predominant in leukemia; arginine R at position 140 is replaced by another amino acid, giving a missense mutation)*

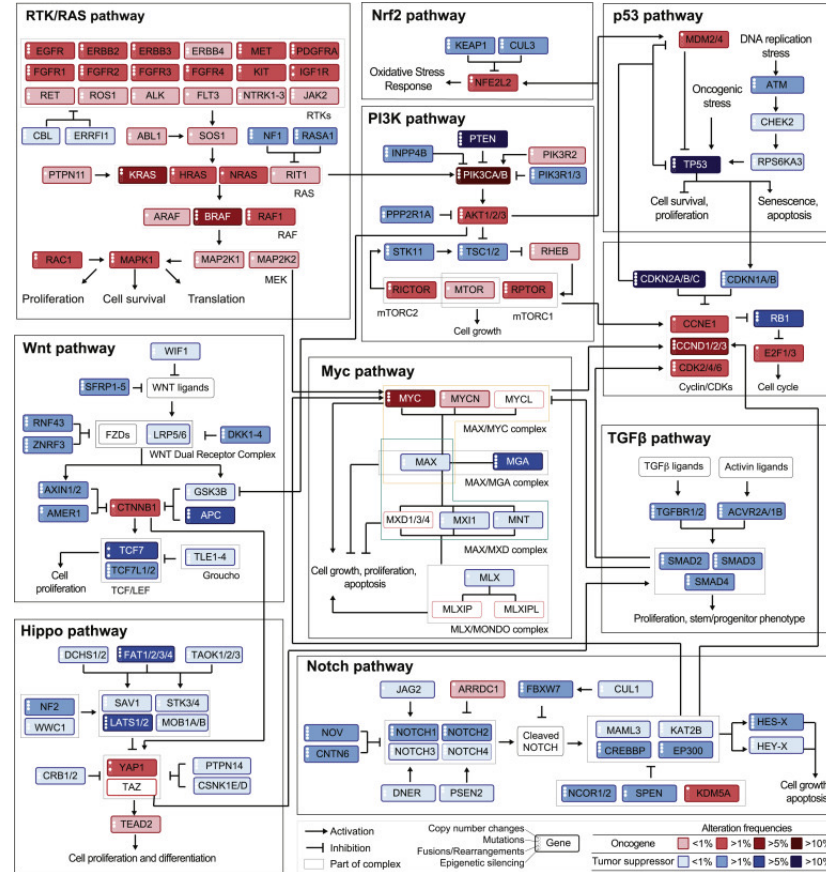
Production of new metabolites: 2-hydroxyglutarate (2-HG)



Production of new metabolites: 2-HG influence DNA methylation



EPFL Many pathways are interconnected (network)



Multiple signals



**Converge
intracellular signaling**



Readout



proliferation



cell survival

Multiple signals



**Converge
intracellular signaling**

Transcription factors



Metabolism



Epigenetics



Readout



**promote
proliferation**

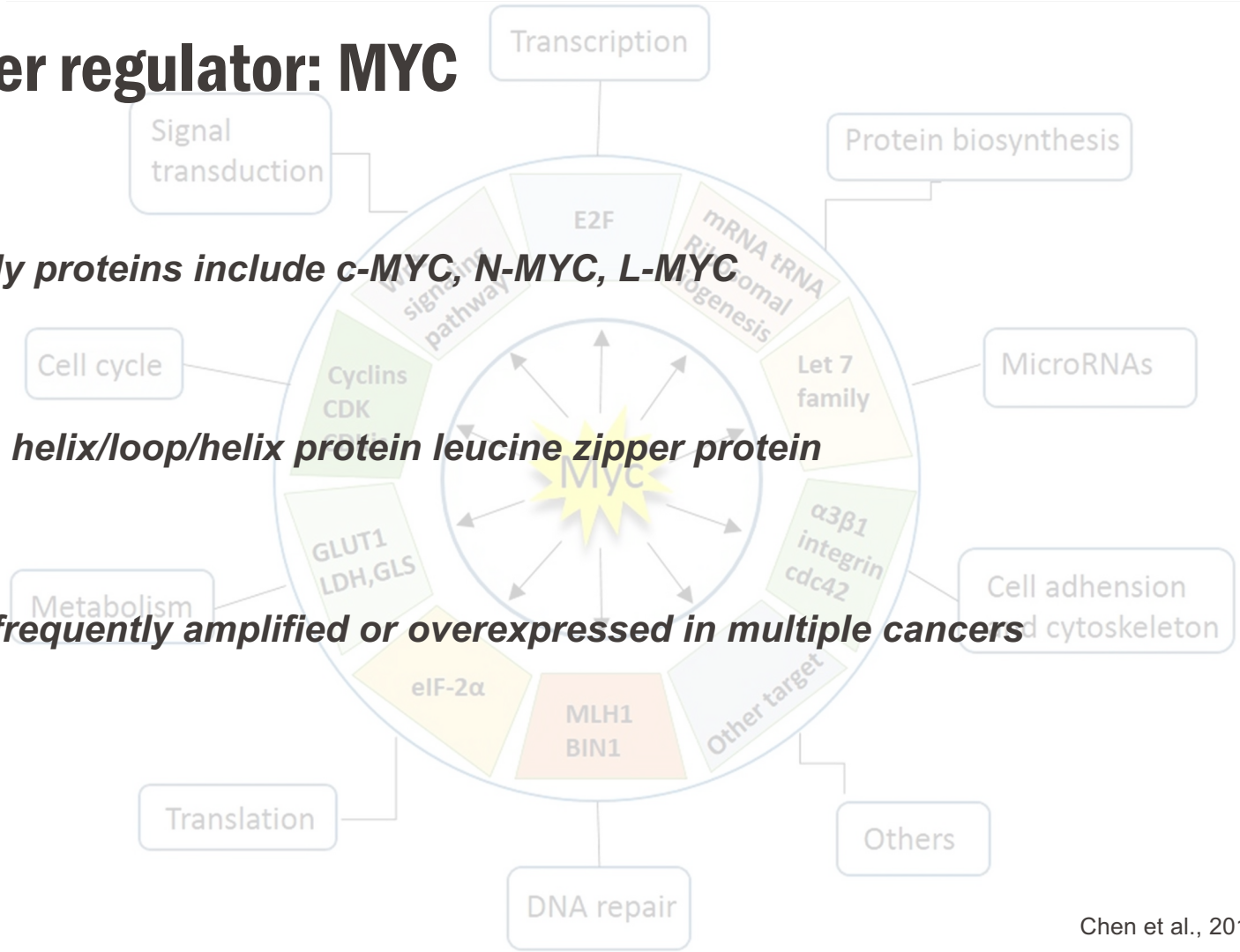
**promote
cell survival**

A master regulator: MYC

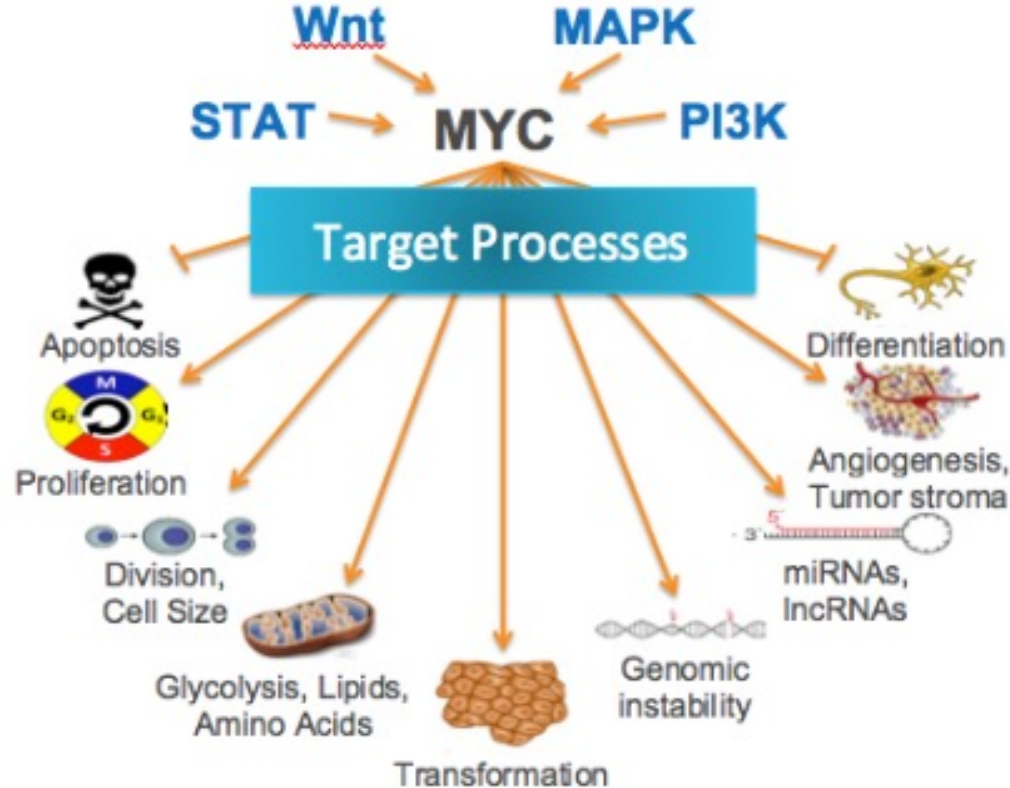
MYC family proteins include c-MYC, N-MYC, L-MYC

MYC is an helix/loop/helix protein leucine zipper protein

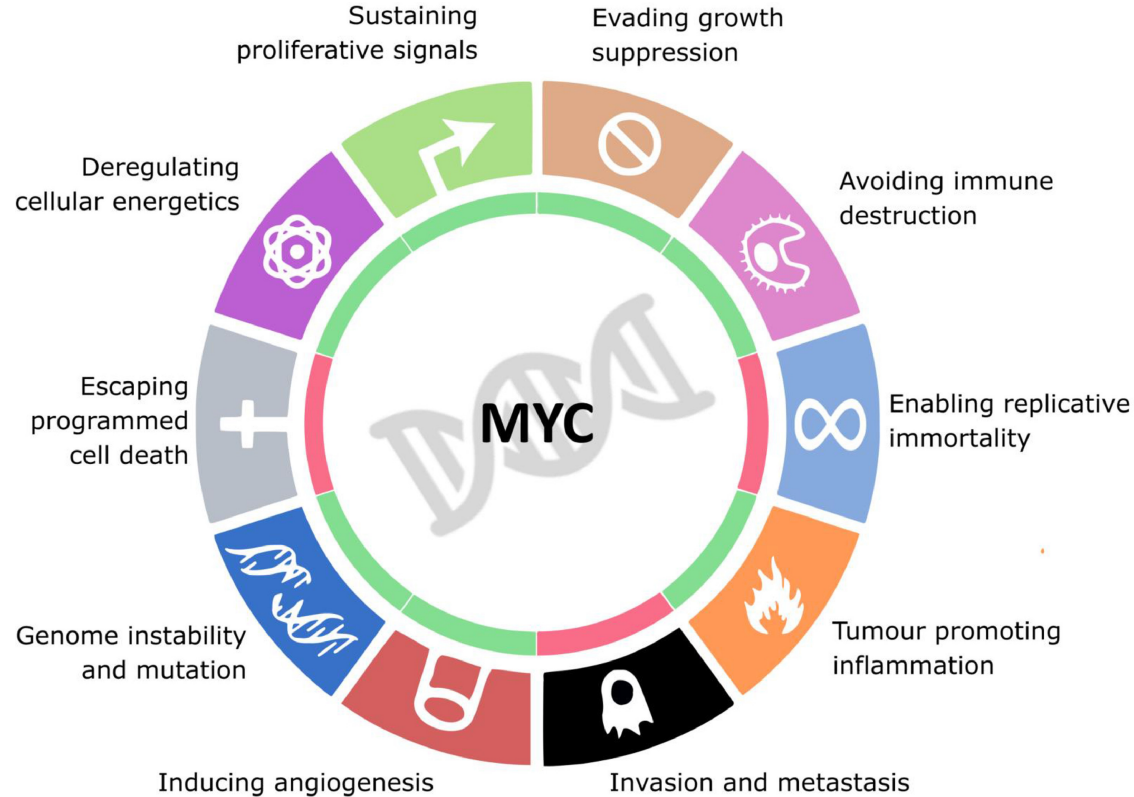
c-MYC is frequently amplified or overexpressed in multiple cancers



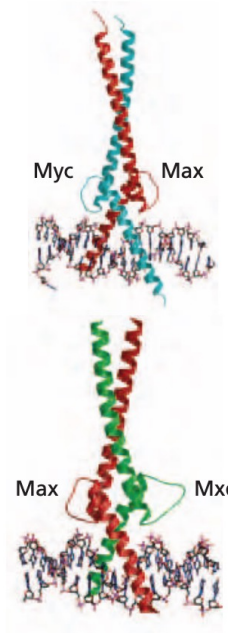
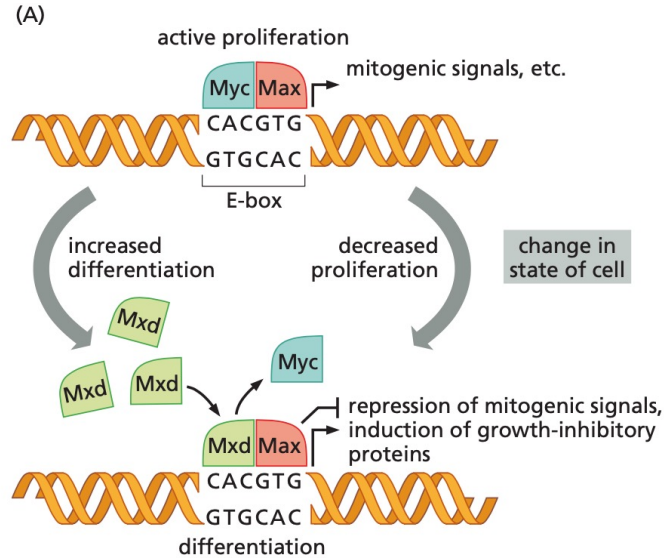
MYC is a transcription factor and potent oncogene



MYC regulates all the hallmarks of cancer



How does MYC control several biological processes?

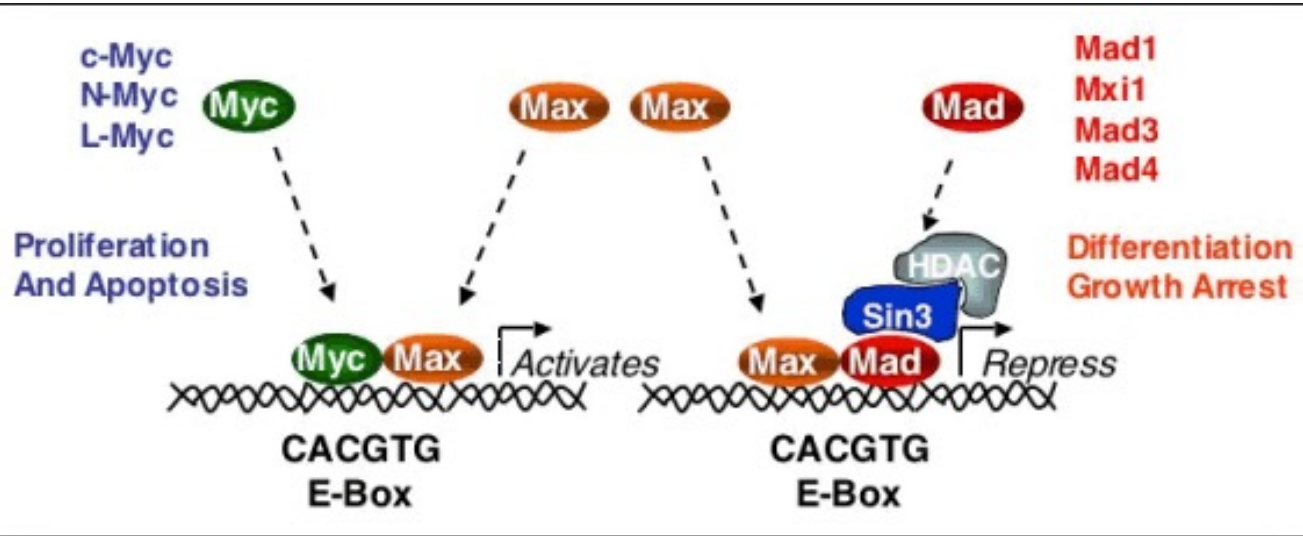


MYC forms dimer with MAX

MYC/MAX complex binds specific DNA sequence (E-BOX)

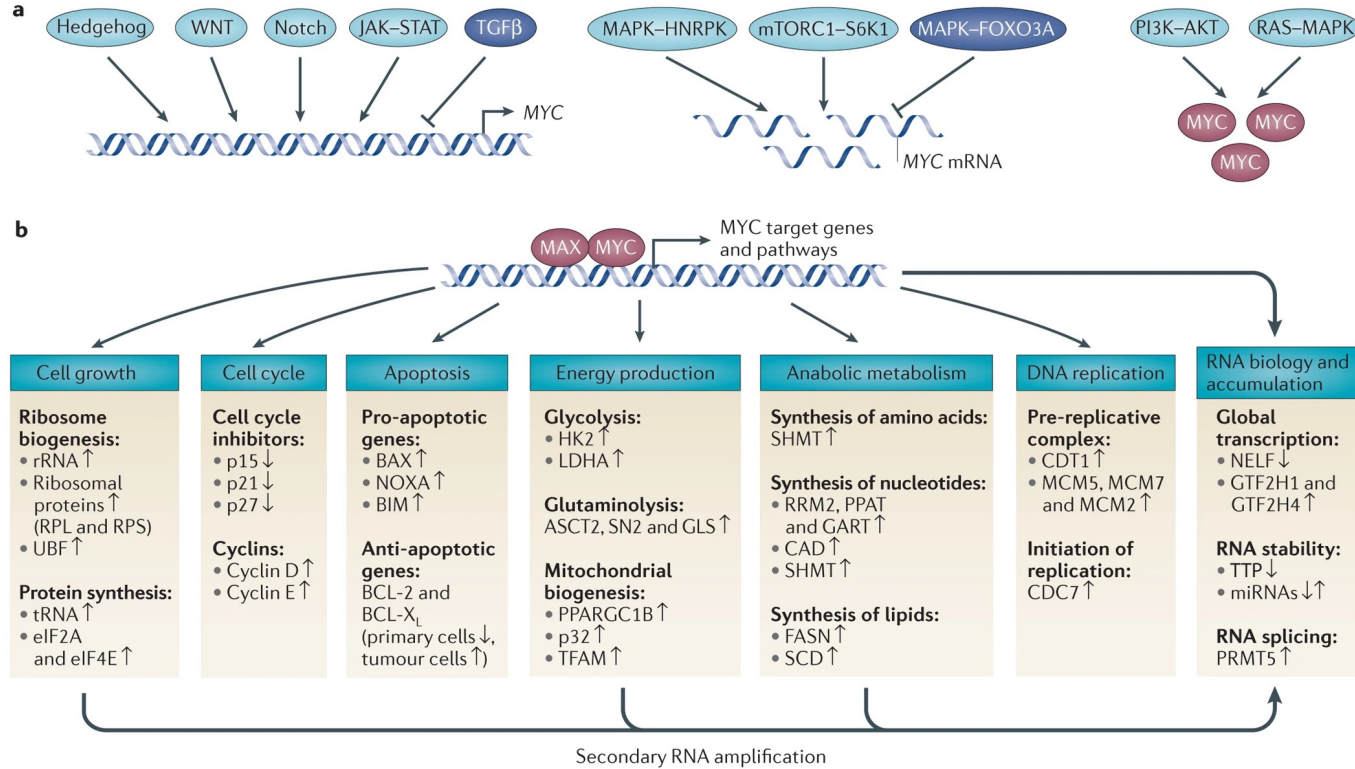
MYC activates the expression of many genes

MYC regulates gene expression



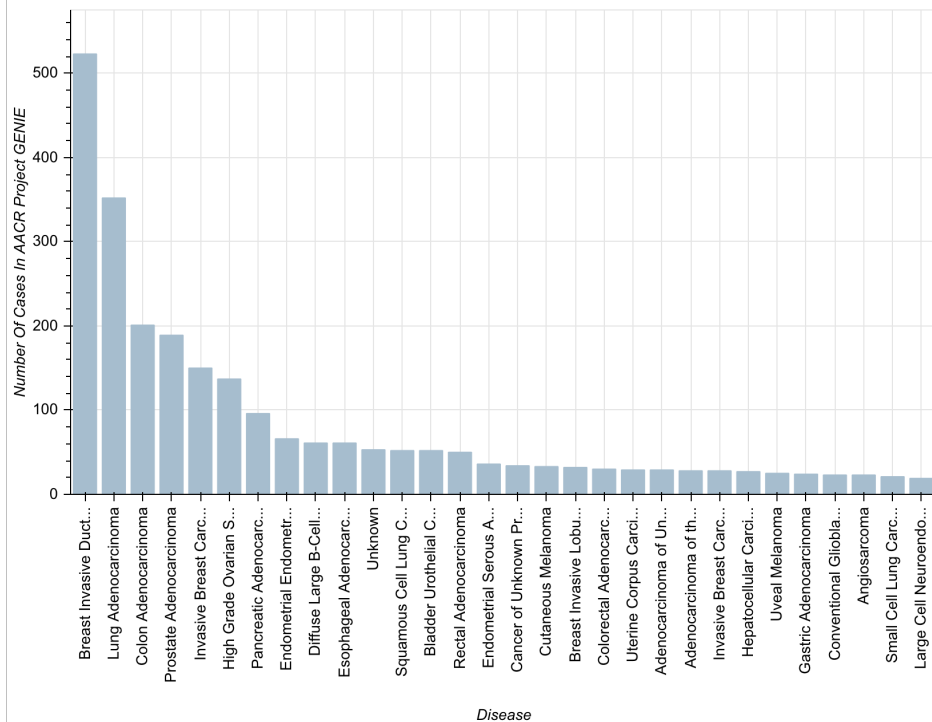
Antagonistic effects of Max partners

MYC targets

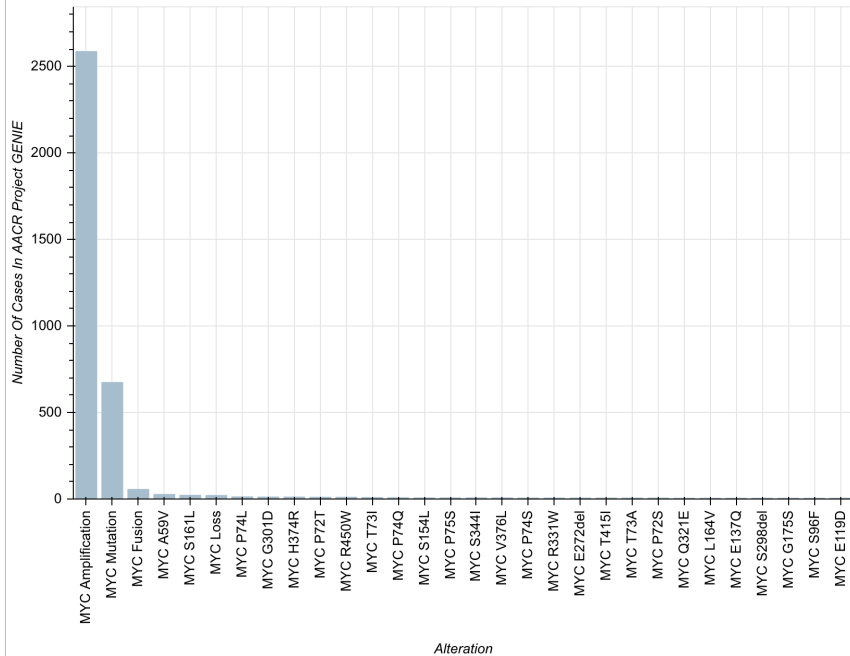


MYC alterations in cancer

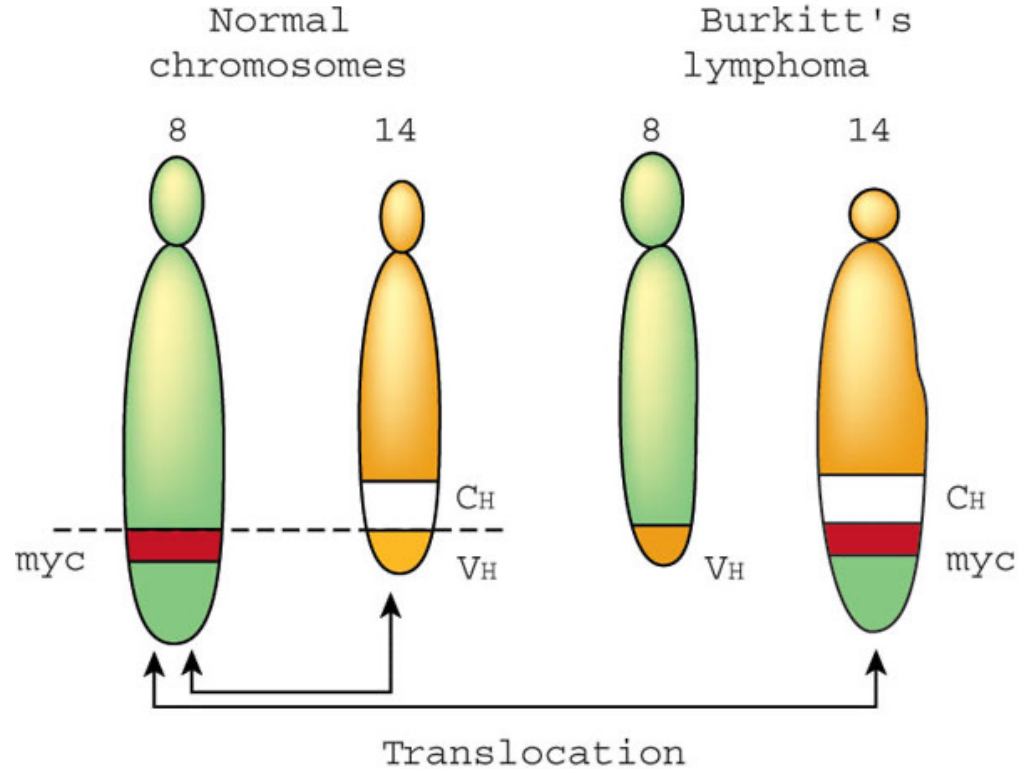
Myc alterations in different malignancies



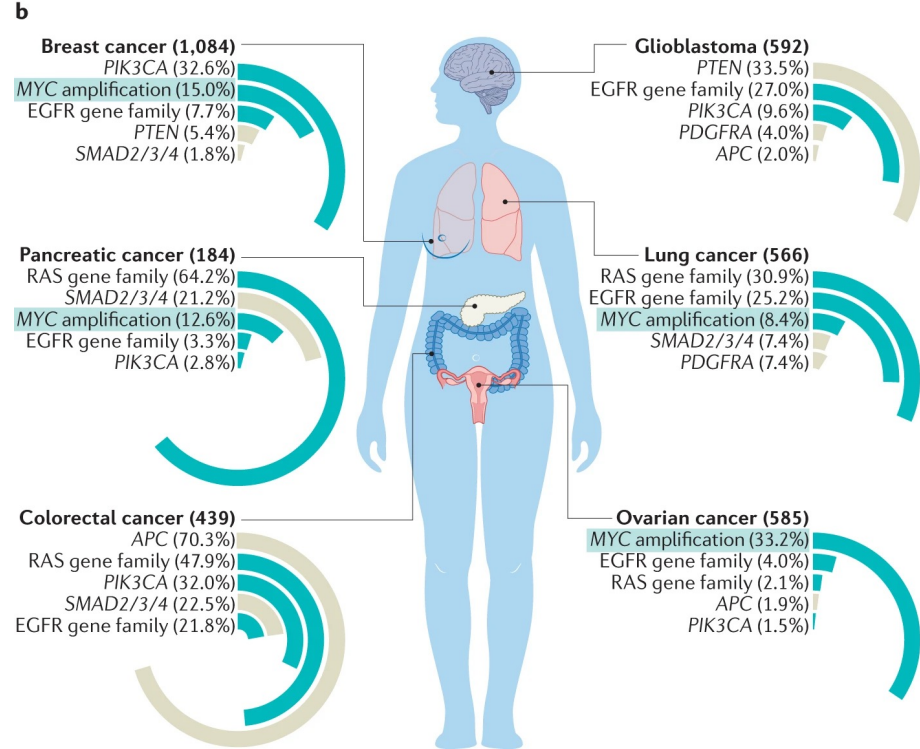
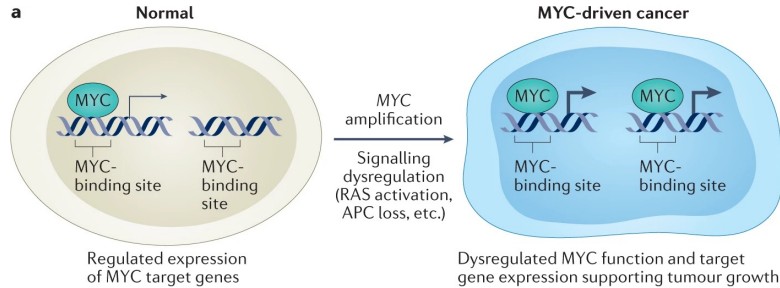
Most common Myc alterations



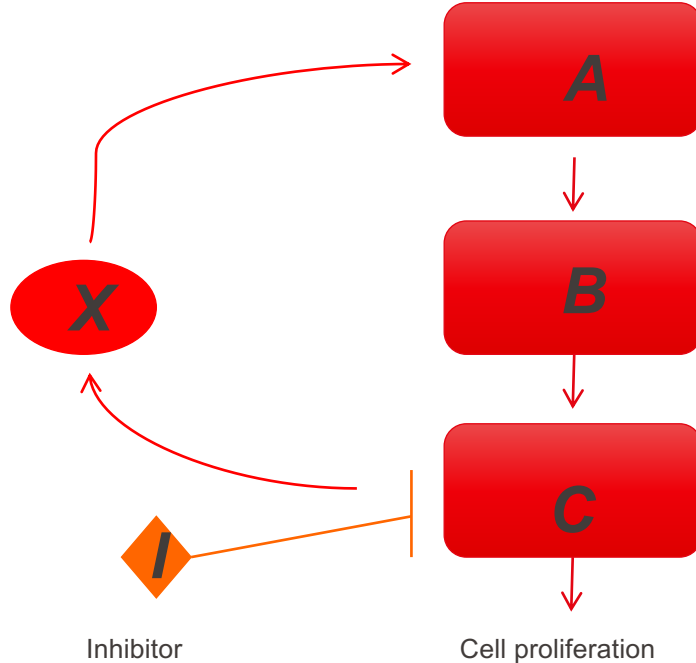
MYC translocation in Burkitt lymphoma



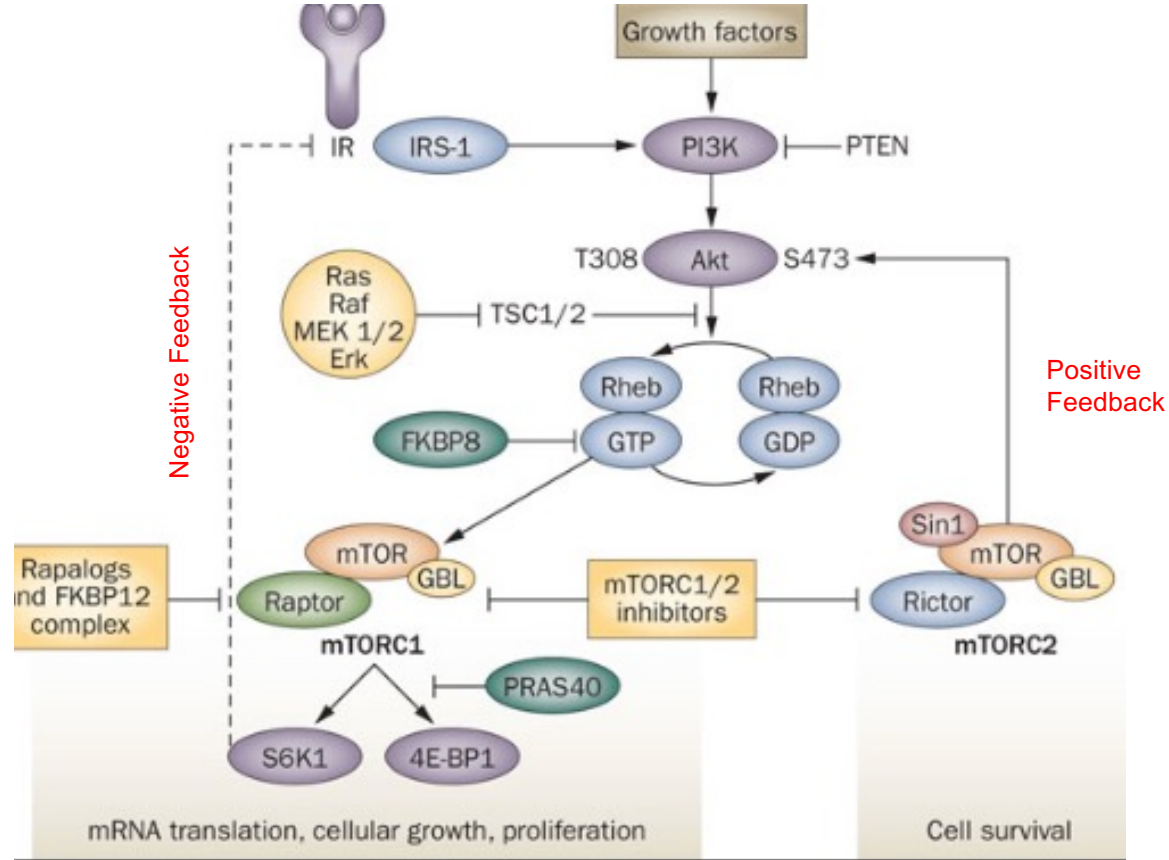
MYC amplification (copy number increase) in cancer



The pathway signal is not linear but can be supported by feedback activations



mTOR re-activates AKT



How can we analyze a pathway's activity?

Visualizing the signaling:

Reporter constructs

Protein-protein interactions (FRET)

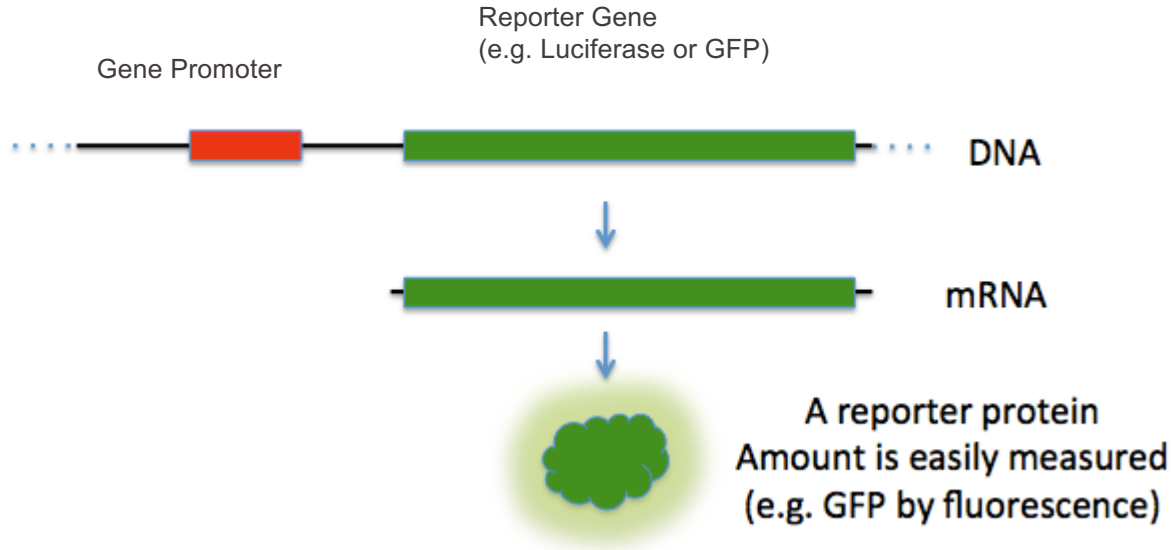
Understand the signaling connection:

yeast two hybrid screen/affinity purification/mass spectrometry

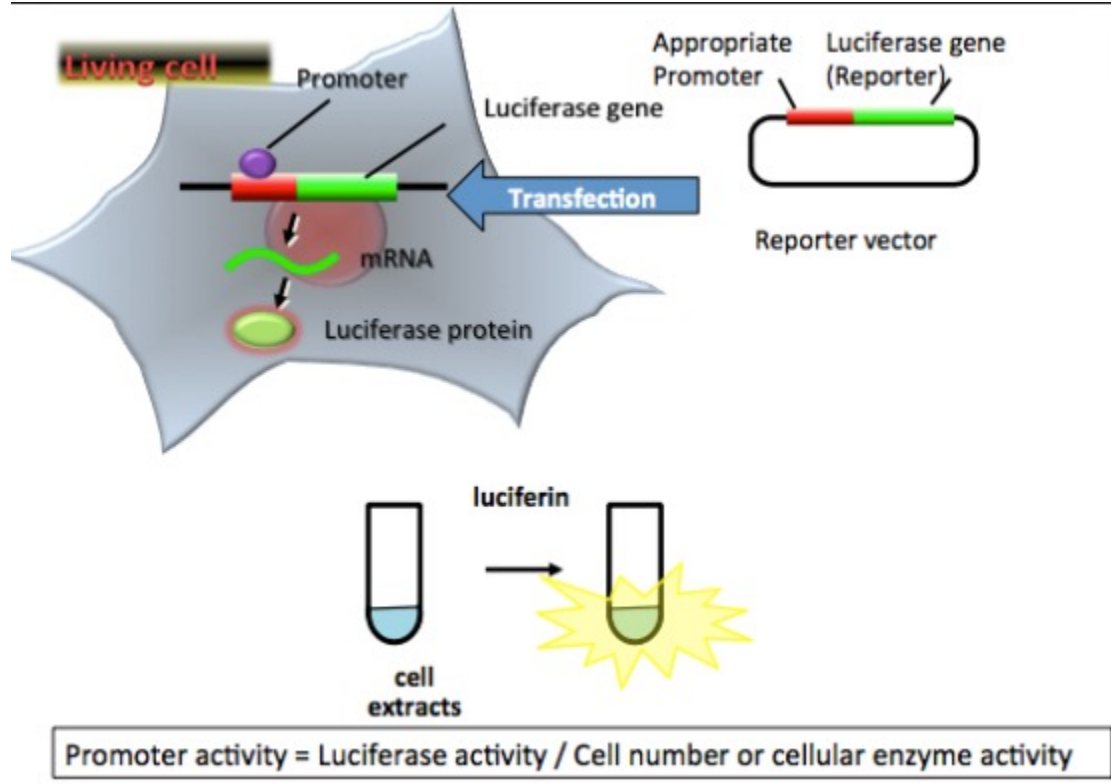
Mass-cytometry (CytoF)

Reporter Construct

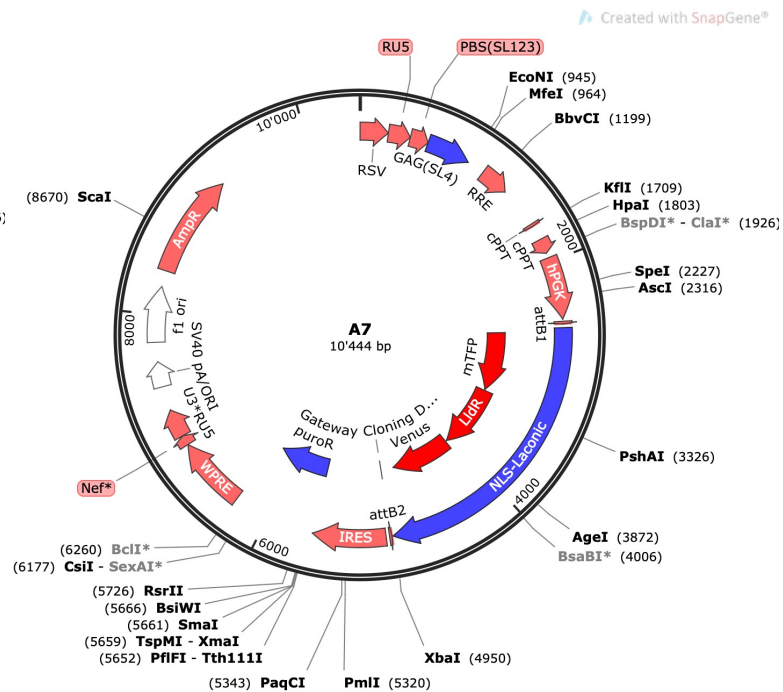
To understand if a transcription factor regulates the expression of specific genes



Reporter Construct



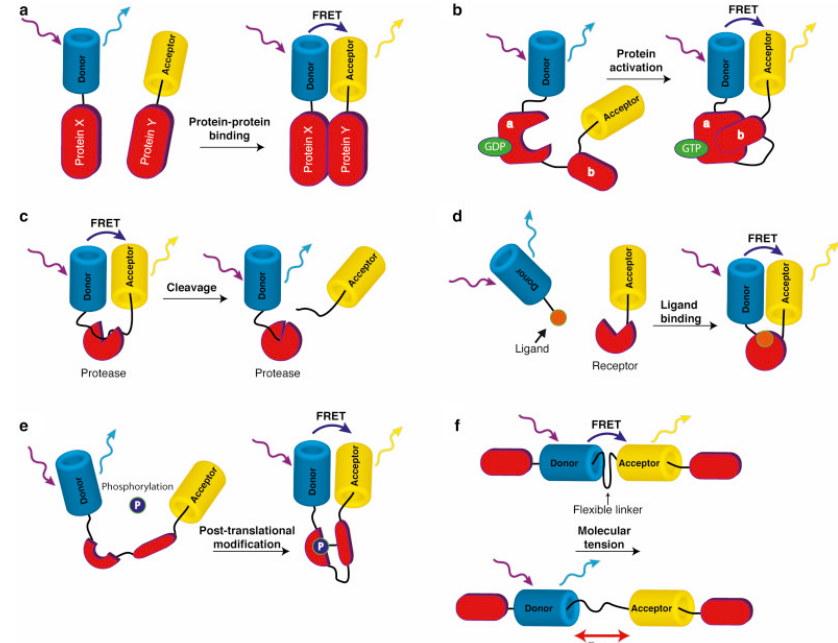
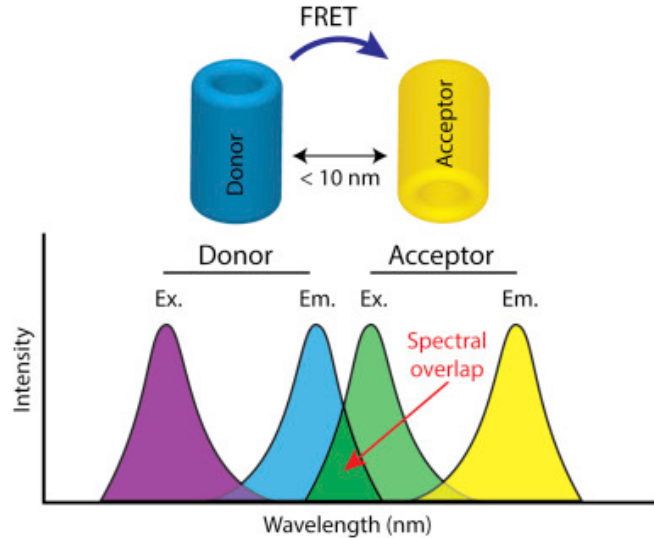
lactate reporter



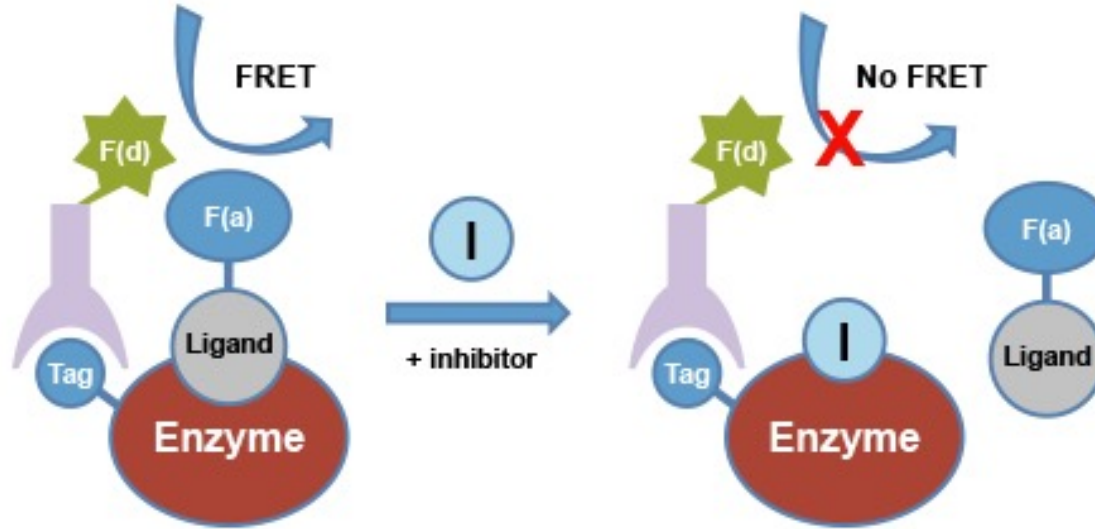
FRET (Förster Resonance Energy Transfer)

Used to study (in living cells):

Protein-protein interactions
Ligand binding to a receptor
Molecular dimerization



FRET with specific inhibitors



How can we analyze the pathways activity?

Visualizing the signaling:

Reporter constructs

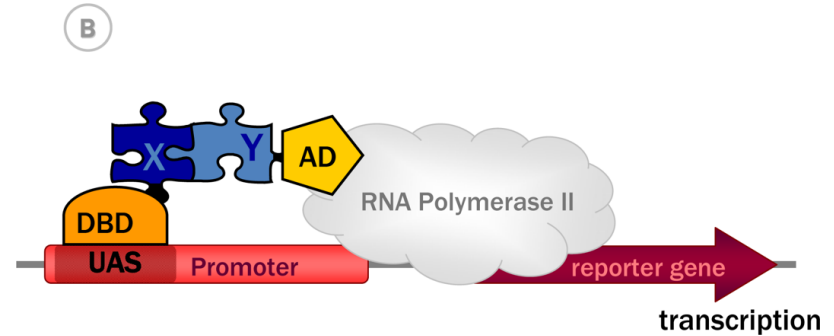
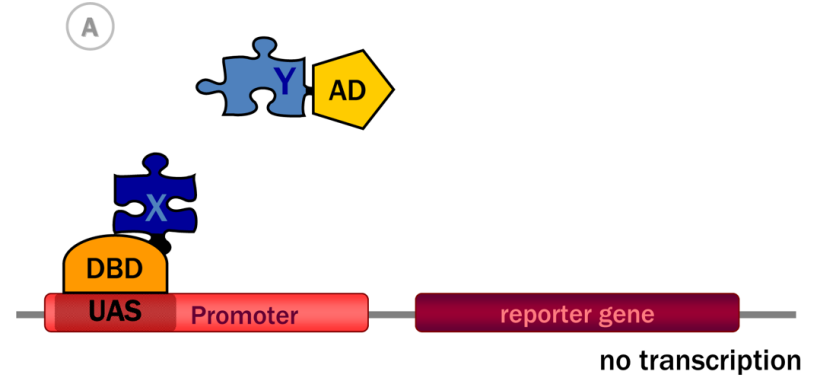
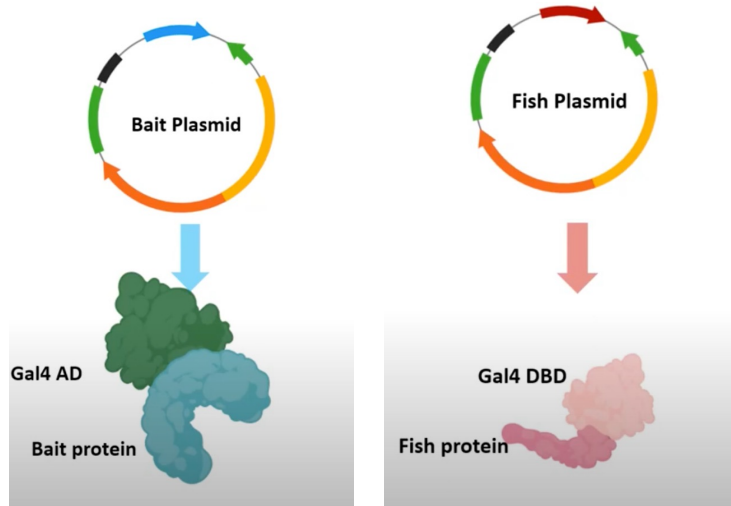
Protein-protein interactions (FRET)

Understand the signaling connection:

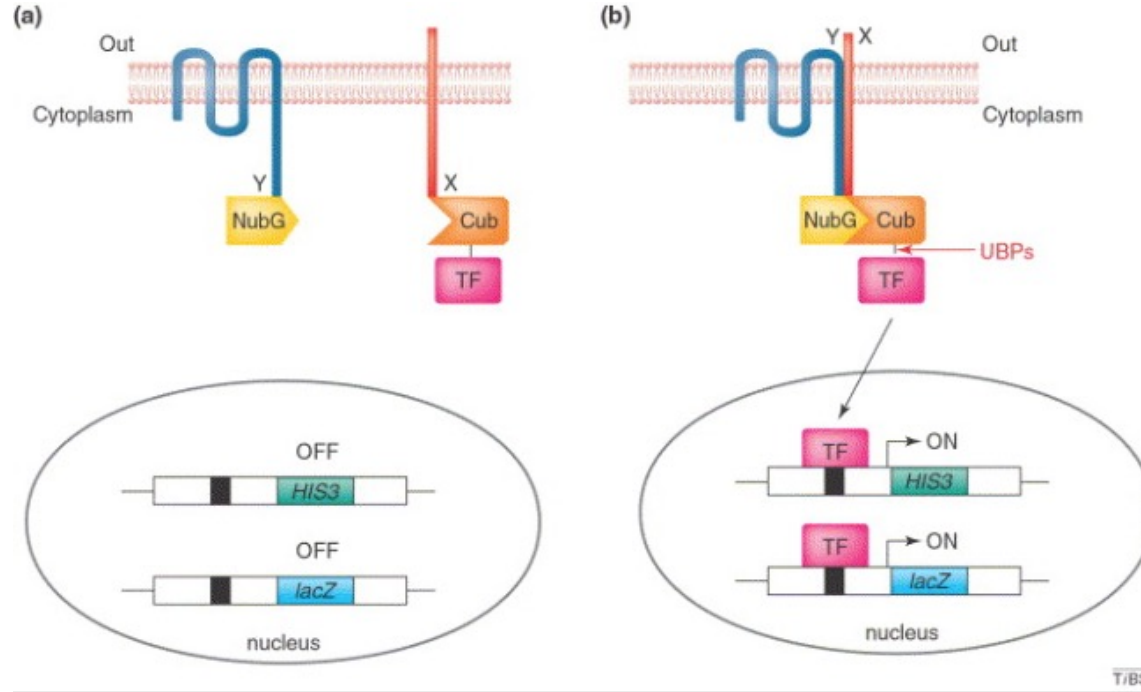
yeast two hybrid screen/affinity purification/mass spectrometry

Mass-cytometry (CyTOF)

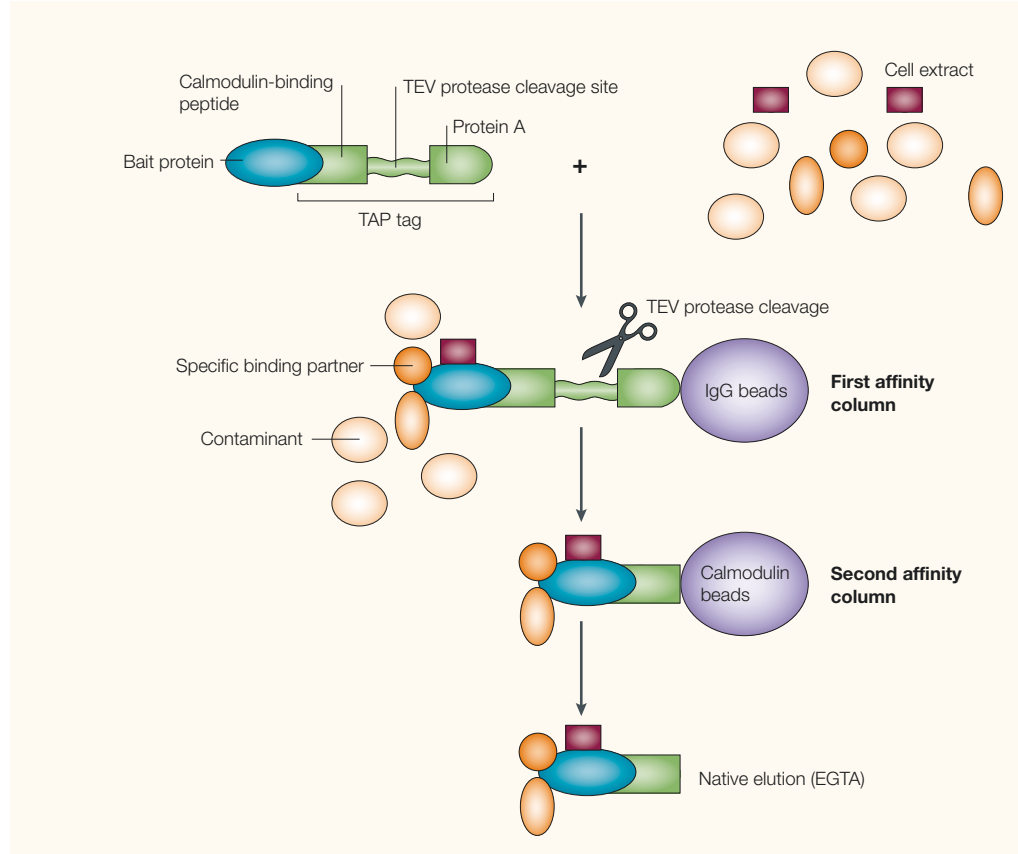
Yeast two hybrid screen (to discover protein-protein interactions)



EPFL Yeast two hybrid screen (transcription activation)



EPFL Protein-protein interaction by affinity purification



Frequently altered pathways in Cancer

<https://www.pnas.org/content/pnas/105/2/692.full.pdf>