

### Cell death or survival:

- > Interactions between signaling pathways
- > Survival factors: classical and alternative pathways

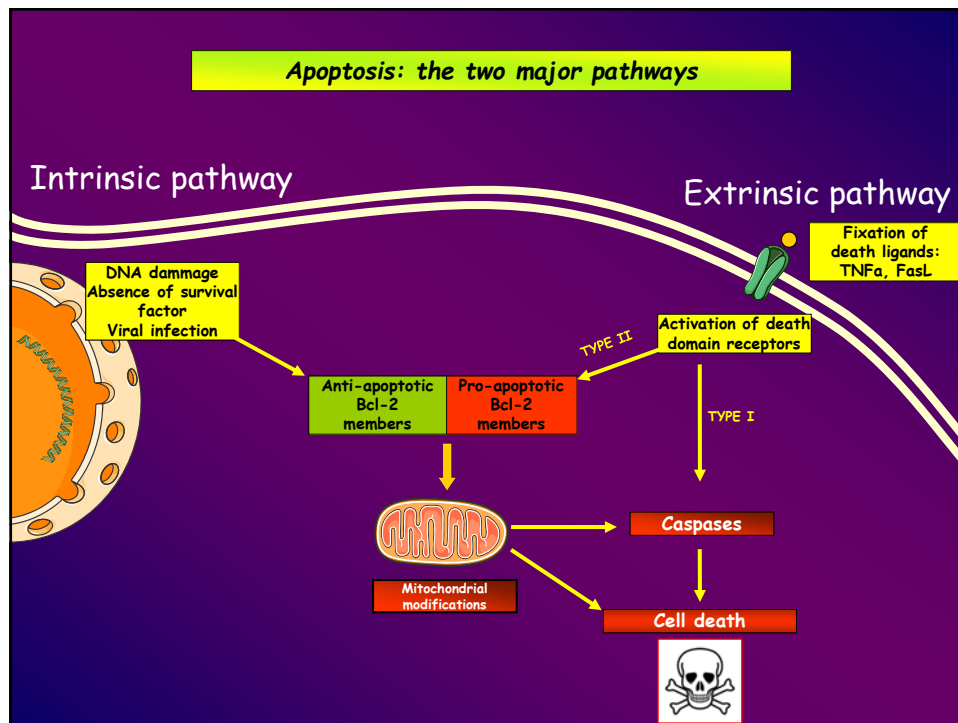
Flore Renaud

Orsay, novembre 2025

UMR 9019 CNRS/U. Paris Saclay/EPHE-PSL,

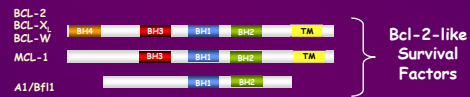
Team « Mechanisms of DNA Repair and Carcinogenesis », Institut Gustave Roussy, Villejuif

flore.renaud-paitra@ephe.psl.eu; flore.renaud-paitra@gustaveroussy.fr

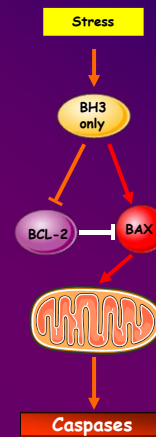
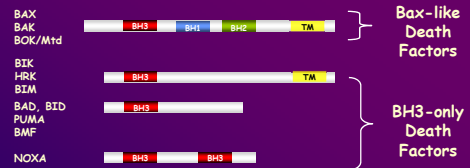


## The BCL-2 family: regulation of apoptosis

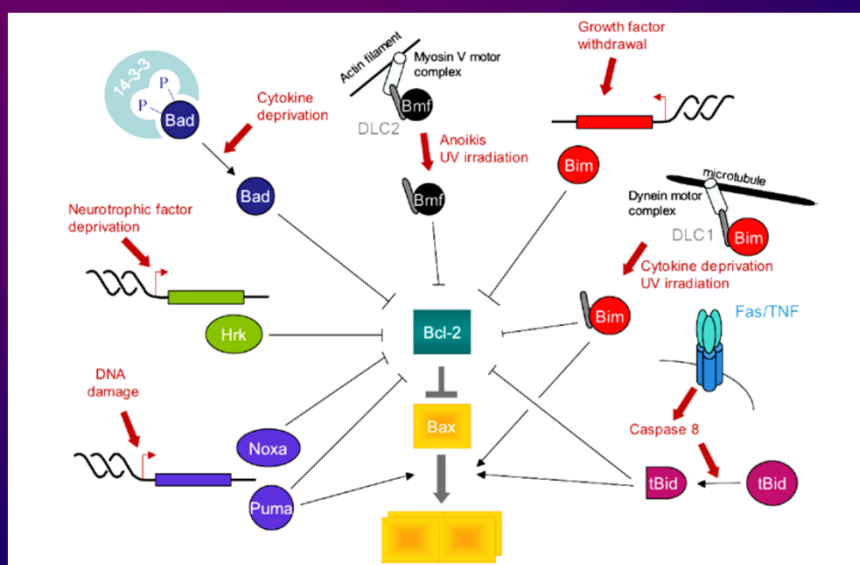
### Anti-apoptotic members



### Pro-apoptotic members

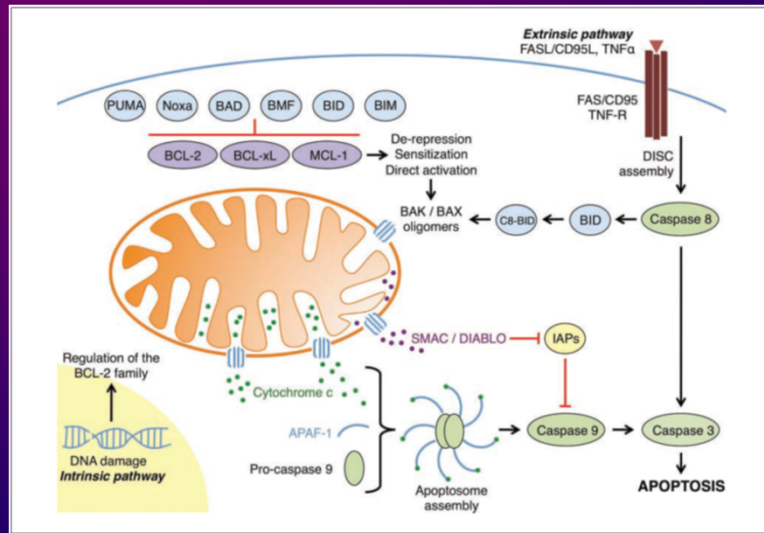


## The BH3-only death factors activation



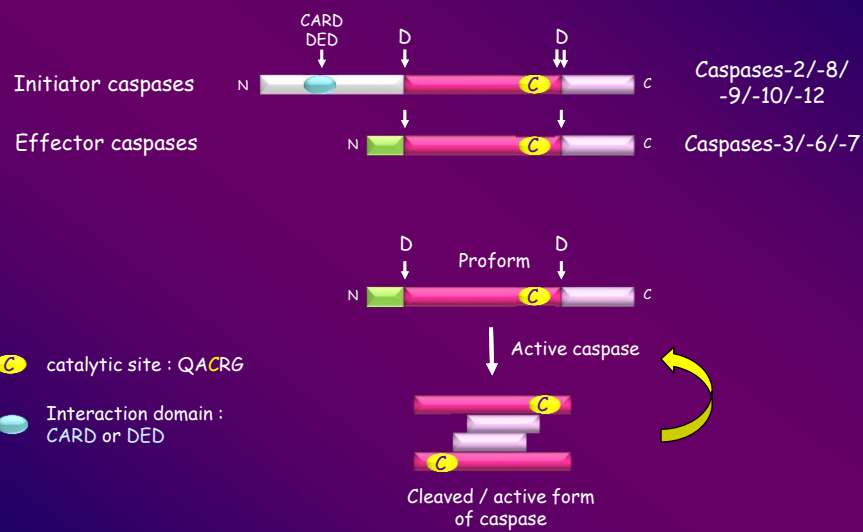
Lucken-Ardjomande and Martinou 2005 C.R. Biologies

### The BCL-2 family: regulation of apoptosis

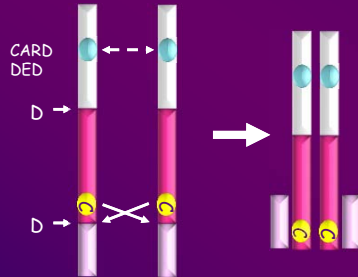


Elkhomi et al. 2011 Genes and Cancer

### The caspases : effectors of cell death



### Activation of initiator caspases

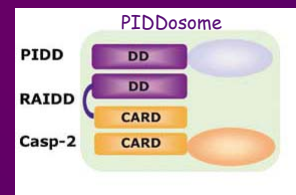
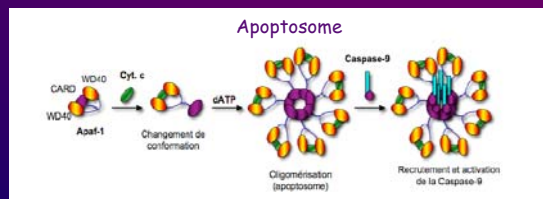
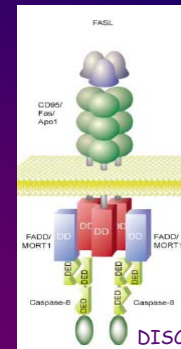


Activation of initiator caspases:

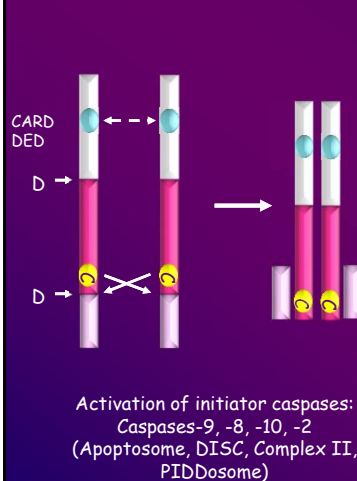
Caspase-9 (CARD): Apoptosome

Caspases-8, -10 (DED): DISC, Complex II

Caspase-2 (CARD): PIDDosome,



### Activation of the caspase cascade



Activation of initiator caspases:  
Caspases-9, -8, -10, -2  
(Apoptosome, DISC, Complex II,  
PIDDosome)

Activation of effector caspases:  
Caspases-3, -6, -7

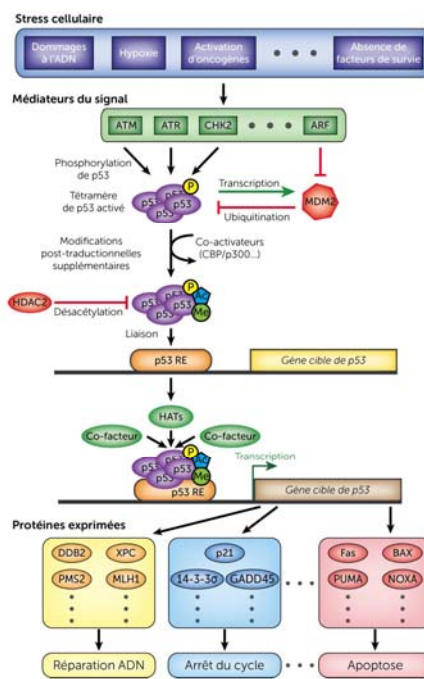
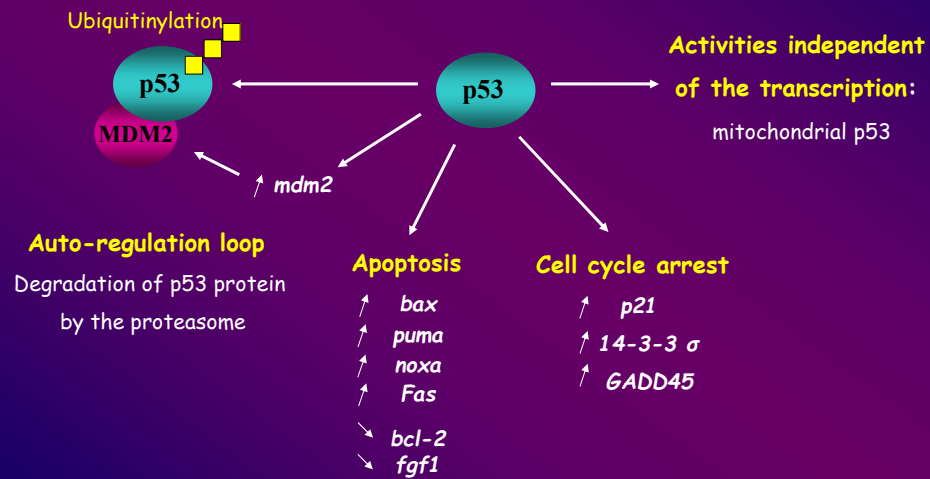
Cleaving many proteins:  
Activation of caspases  
(amplification)

Activation of proteins  
involved in apoptosis  
(ICAD, PARP, Bid, ...)

Inactivation of proteins  
required for cell  
survival (Bcl-2, Bcl-X<sub>L</sub>,  
XIAP, MDM2, Rb, CREB,  
EGF-R, MEK, Raf-1, ...)

Degradation of  
cytoskeletal proteins  
(lamin, tubulin, actin, ...)

**The oncosuppressor p53: a key regulator of apoptosis**  
**p53 is a transcription factor involved in**  
**apoptosis and cell cycle arrest**

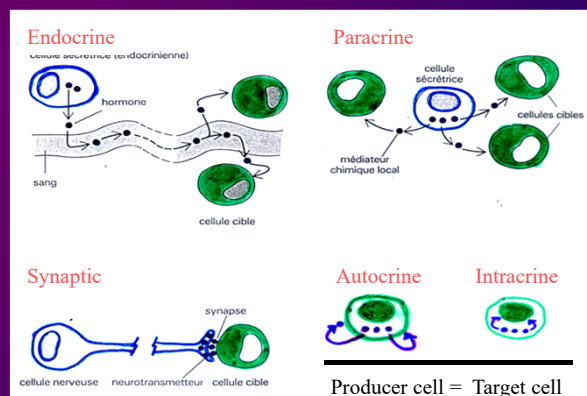


Caroline Pirou  
Thèse 2016

### Growth factor families

- EGF (Epidermal Growth Factor)
- PDGF (Platelet-derived growth factor)
- Growth Hormones
- VEGF (Vascular endothelial growth factor)
- FGF 1-23 (Fibroblast Growth Factor)
- Neurotrophins: NGF, NT3, NT4, BDNF
- GDNF (Glial cell line-derived neurotrophic factor), NTN (neurturin), PSP (persephin)
- Neurokines: CNTF, LIF
- Insulin, IGF-I, IGF-II

### The growth factor pathways



**The paracrine pathway:** the growth factor is produced and secreted by a cell and acts on neighboring cells.

**The endocrine pathway:** the growth factor is produced and secreted by a cell and then passes through the blood or lymphatic circulation to act on cells which may be very distant (Insulin, IGF-I/II)

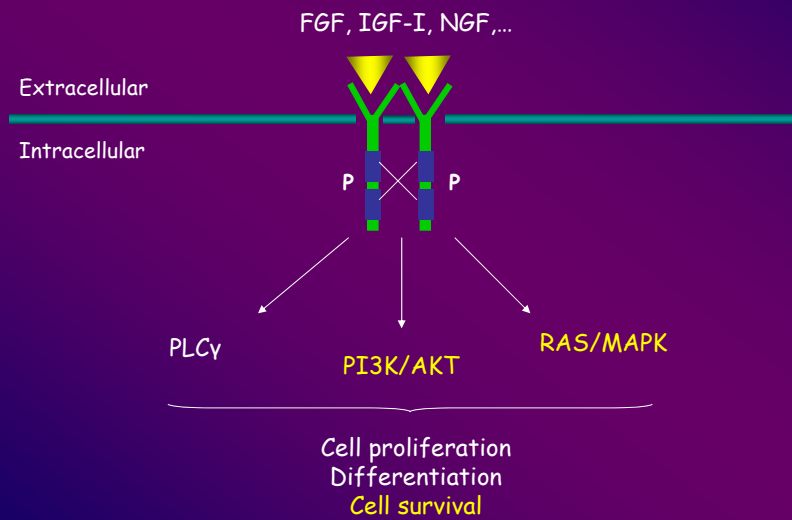
**The synaptic pathway:** the growth factor is produced by a neuronal cell and then transported along the axon to synapses for acting on cells which may be very distant (neurotrophic factors)

For **autocrine and intracrine pathway**, the producer cell is the target cell.

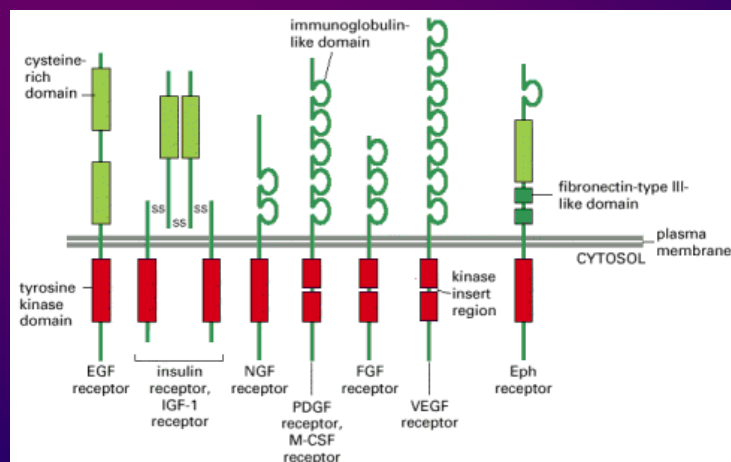
**The autocrine pathway:** the cell produces the factor and its receptors and therefore the factor is produced and secreted to act on the producer cell.

**The intracrine pathway:** the cell produces the factor but it is not secreted and it acts directly into the cell by an internal pathway independent of the receptors.

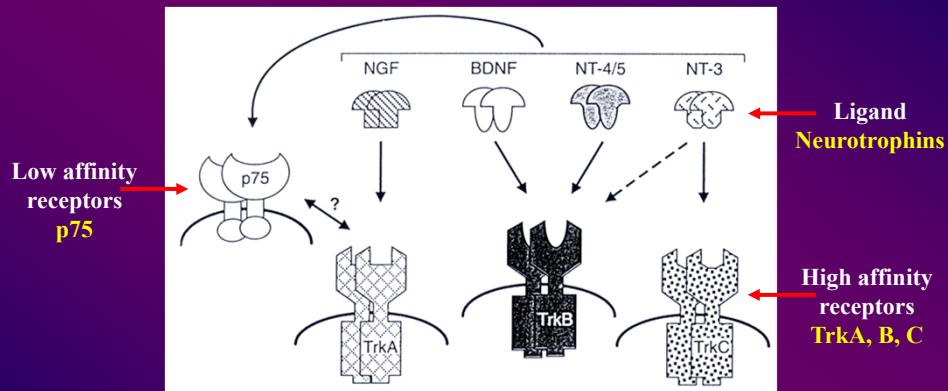
*The classical pathway of action of growth/survival factors involves activation of plasma membrane receptor tyrosine kinases (paracrine, autocrine, endocrine and synaptic)*



### *The receptor tyrosine kinases (RTK)*

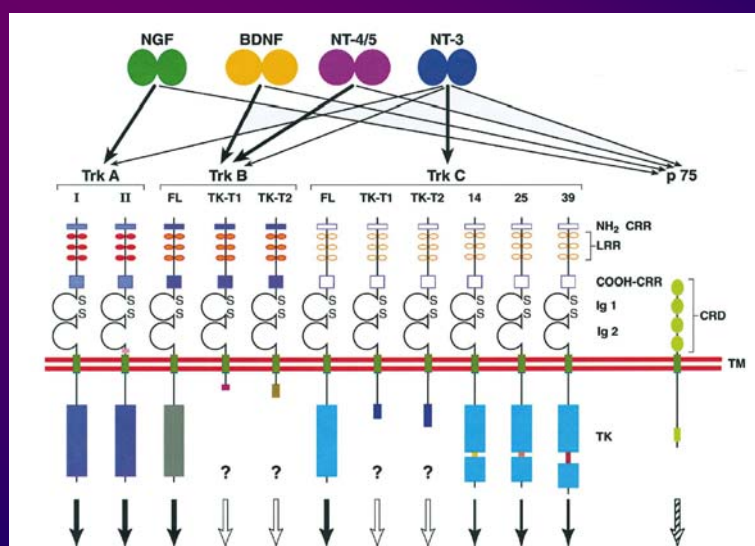


### The High affinity receptors (RTKs) and Low affinity receptors



In addition to high affinity receptors (RTK), there are low-affinity receptors, in particular for the neurotrophic factors. The low affinity receptors either increase growth factor stability and/or increase high affinity receptor activity. They could also induce RTK independent pathways.

### The neurotrophin receptors (TrkA/B/C and p75)





### The two major signaling pathways of the neurotrophic/survival factors

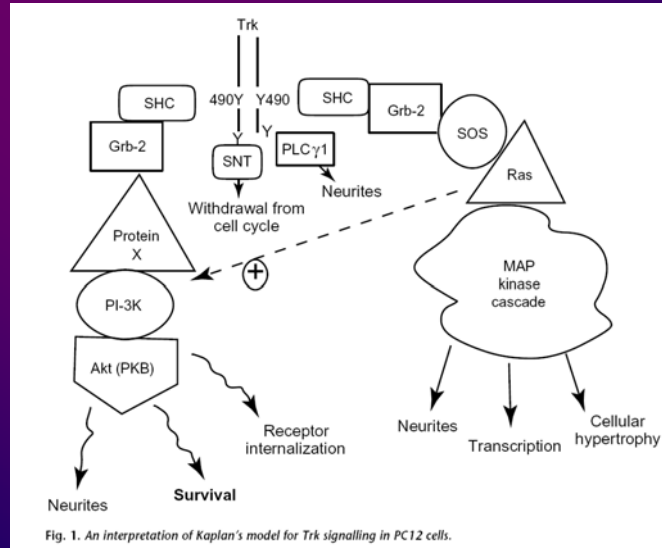
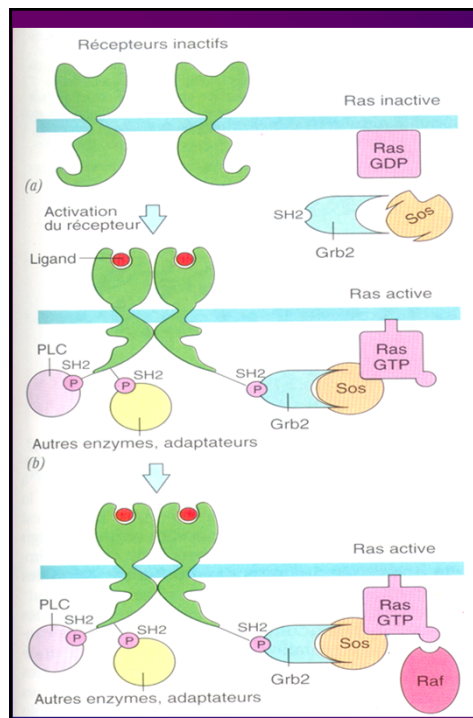


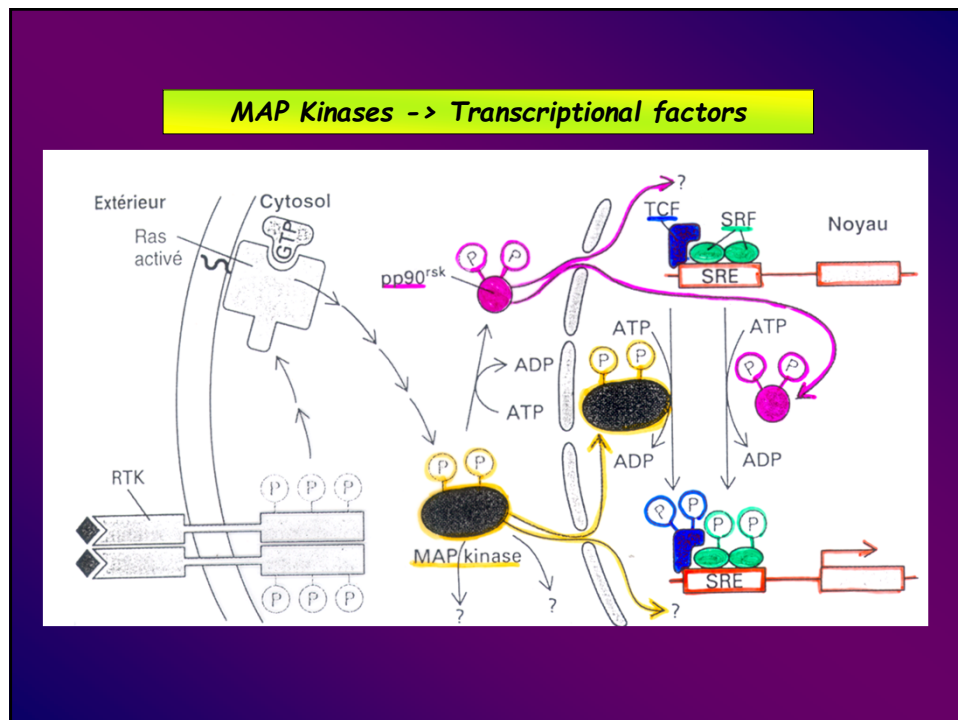
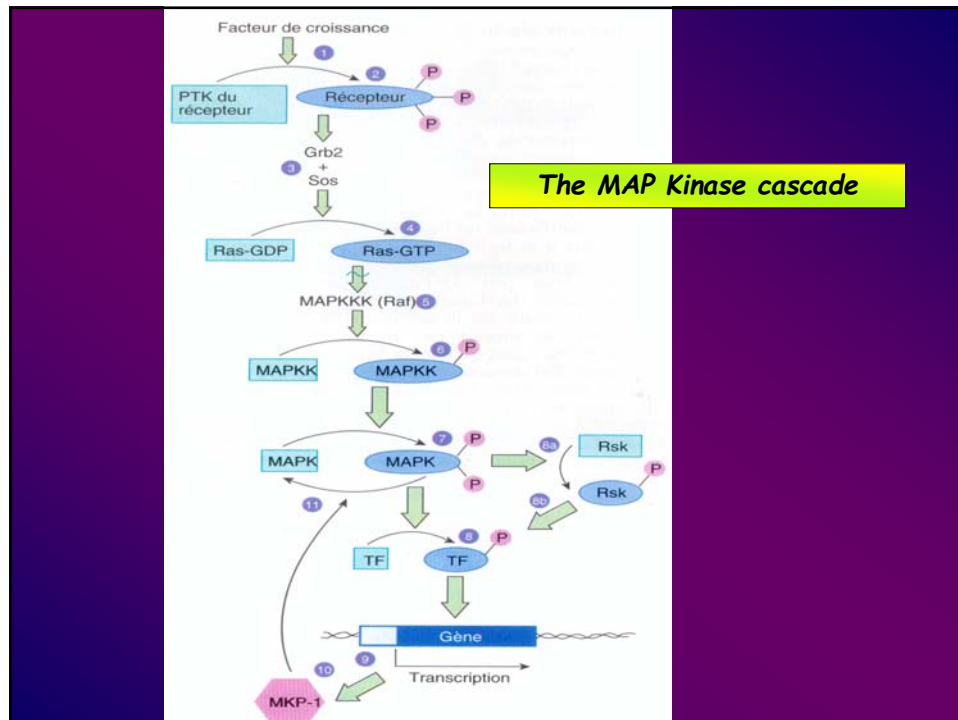
Fig. 1. An interpretation of Kaplan's model for Trk signalling in PC12 cells.

Tolkovsky 1997 TINS

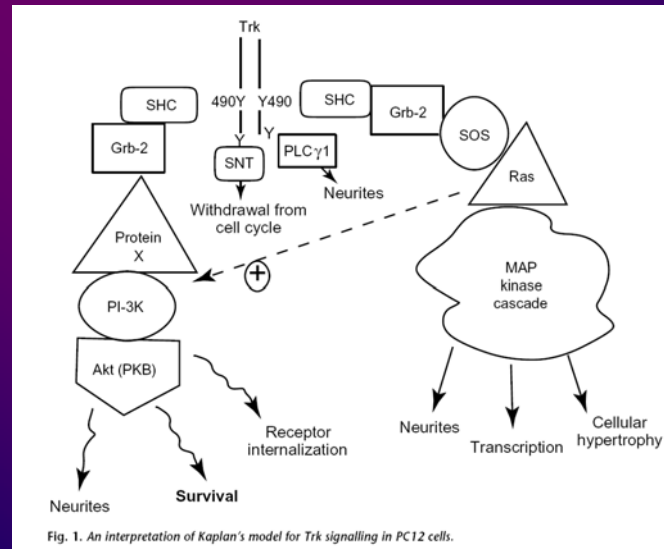


### Activation of Ras by RTKs

- In the absence of growth factor, the receptors are inactive monomers. Ras is combined with GDP and therefore inactive.
- Ligand binding triggers the dimerization and autophosphorylation of the receptors. Phosphotyrosine residues newly formed function as binding sites for Grb2-Sos which induces exchange GDP and GTP on Ras.
- Activated Ras becomes a binding site for Raf and triggers the MAP kinase cascade.



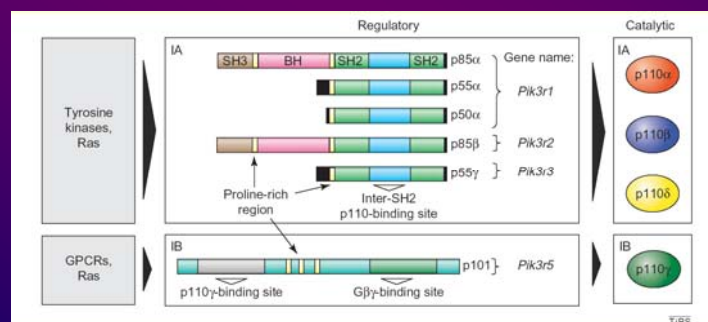
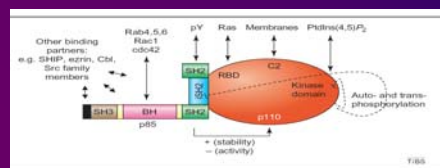
### The two major signaling pathways of the neurotrophic/survival factors



Tolkovsky 1997 TINS

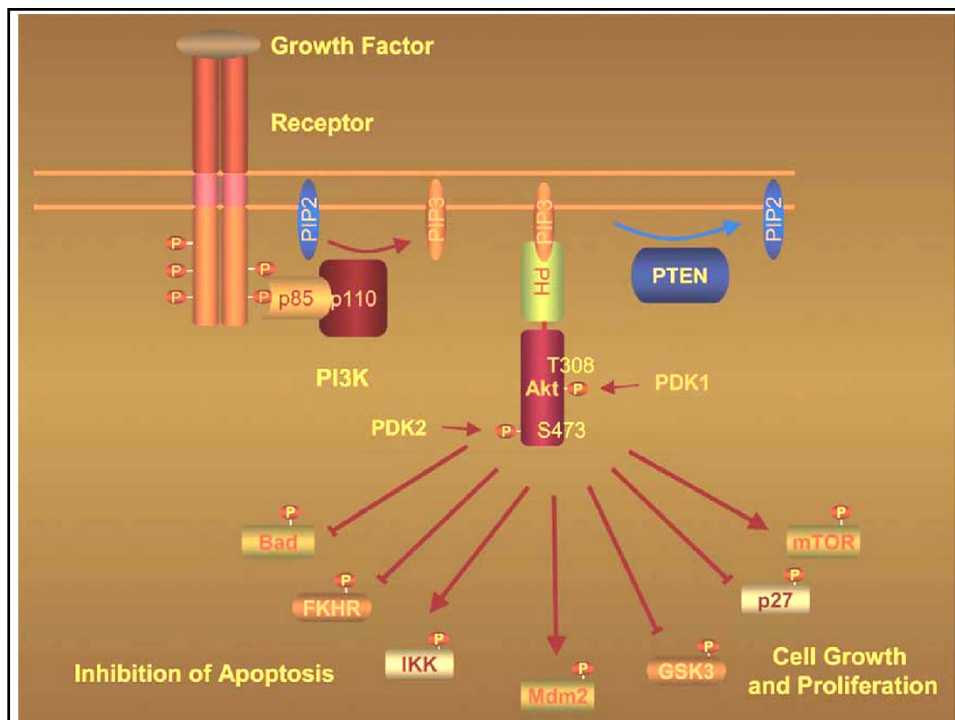
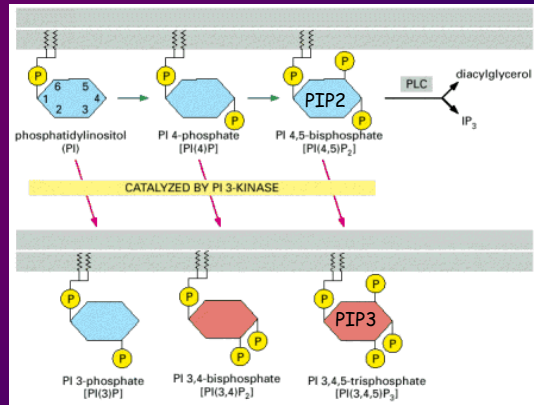
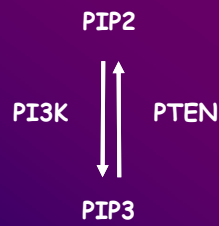
### The PI3K / AKT pathway : the main signaling pathway involved in cell survival

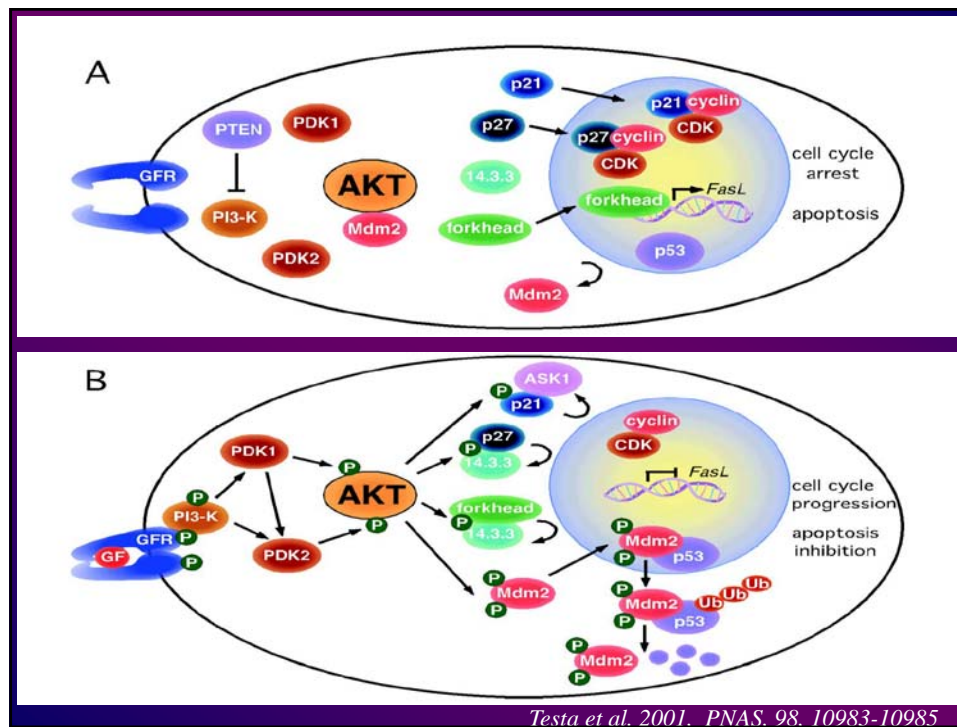
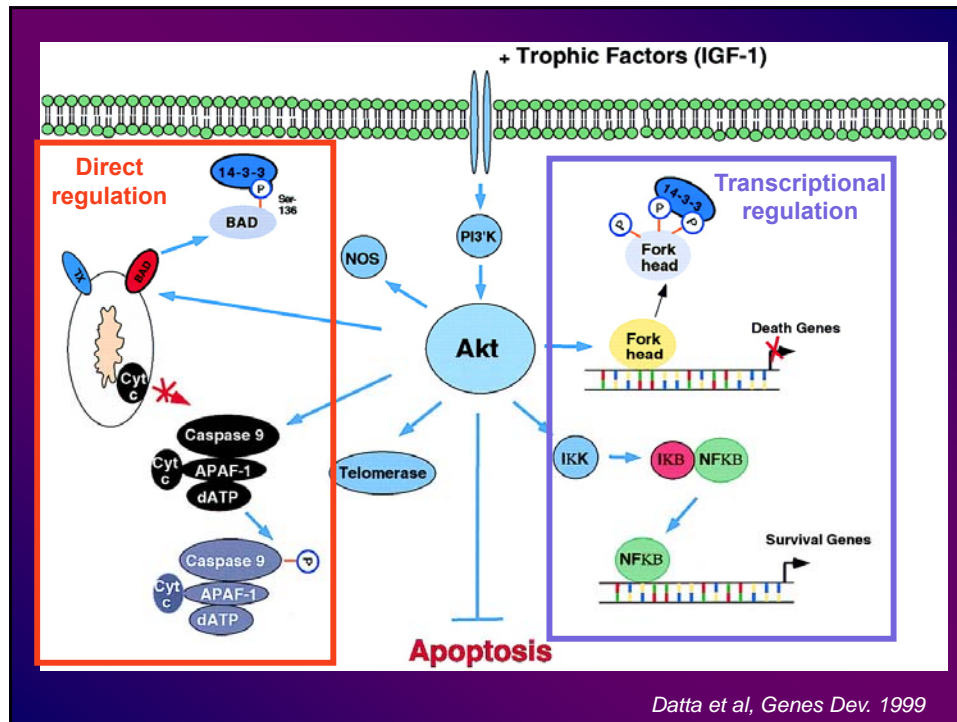
#### The PhosphoInositide 3 Kinase (PI3K)



Vanhaesebroeck et al. 2005, Trends in Biochemical Sciences

**The PI3K / AKT pathway :  
the main signaling pathway involved in cell survival**

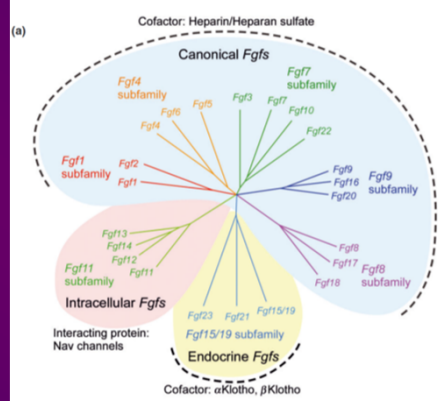




## The fibroblast growth factor signaling pathway

**TABLE 1 | Nomenclature of the Mammalian Fgf and Fgfr family**

Accession Symbol	Name	Alternative Symbol	Name, Comments
<b>(a) Fgf</b>			
FGF1/Fgf1	Fibroblast Growth Factor 1	Abfgf	Acidic Fgf
		Hbgf1	Heparin-binding growth factor 1
		Bfgf	Endothelial cell growth factor
FGF2/Fgf2	Fibroblast Growth Factor 2	Bfgf	Basic Fgf
		Hbgf2	Heparin-binding growth factor 2
FGF3/Fgf3	Fibroblast Growth Factor 3	Wnt-2	Wnt-2 oncogene
		MMTV	MMTV integration site 2
FGF4/Fgf4	Fibroblast Growth Factor 4	Hst1	Human stomach tumor oncogene
		Hst1	Heparin secretory transforming protein 1
		K-Fgf, K-Fgf	Kaposi sarcoma Fgf
FGF5/Fgf5	Fibroblast Growth Factor 5	Hst2	Hst2 oncogene
FGF6/Fgf6	Fibroblast Growth Factor 6	K-Fgf	Keratinocyte growth factor
FGF7/Fgf7	Fibroblast Growth Factor 7	Abfgf	Abrogates induced growth factor
FGF8/Fgf8	Fibroblast Growth Factor 8	K-Fgf	Glia activating factor
FGF9/Fgf9	Fibroblast Growth Factor 9	Elk	Elbow knee synostosis
FGF10/Fgf10	Fibroblast Growth Factor 10	K-Fgf	Keratinocyte growth factor 2
FGF11/Fgf11	Fibroblast Growth Factor 11	Hst3	Fibroblast Growth Factor homologous factor 1
FGF12/Fgf12	Fibroblast Growth Factor 12	Hst4	Fibroblast Growth Factor homologous factor 2
FGF13/Fgf13	Fibroblast Growth Factor 13	Hst5	Fibroblast Growth Factor homologous factor 3
FGF14/Fgf14	Fibroblast Growth Factor 14	Hst6	Fibroblast Growth Factor homologous factor 4
		Spontaneous disease 27	
Fgf15	Fibroblast Growth Factor 15	Spontaneous disease 27	Induced outflow of vertebrate Fgf19
FGF16/Fgf16	Fibroblast Growth Factor 16		Called FGF-13 in some older literature
FGF17/Fgf17	Fibroblast Growth Factor 17		
FGF18/Fgf18	Fibroblast Growth Factor 18		
FGF19	Fibroblast Growth Factor 19		Human outflow of rodent Fgf15
FGF20/Fgf20	Fibroblast Growth Factor 20		
FGF21/Fgf21	Fibroblast Growth Factor 21		
FGF22/Fgf22	Fibroblast Growth Factor 22		
FGF23/Fgf23	Fibroblast Growth Factor 23		

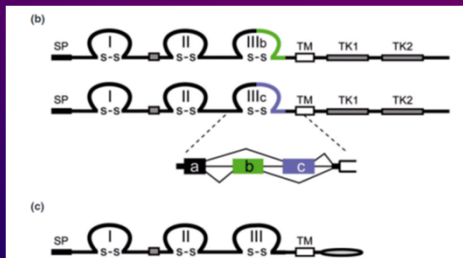


Ornitz and Itoh, Dev. Biol. 2015

## The fibroblast growth factor signaling pathway

**(b) Fgfr**

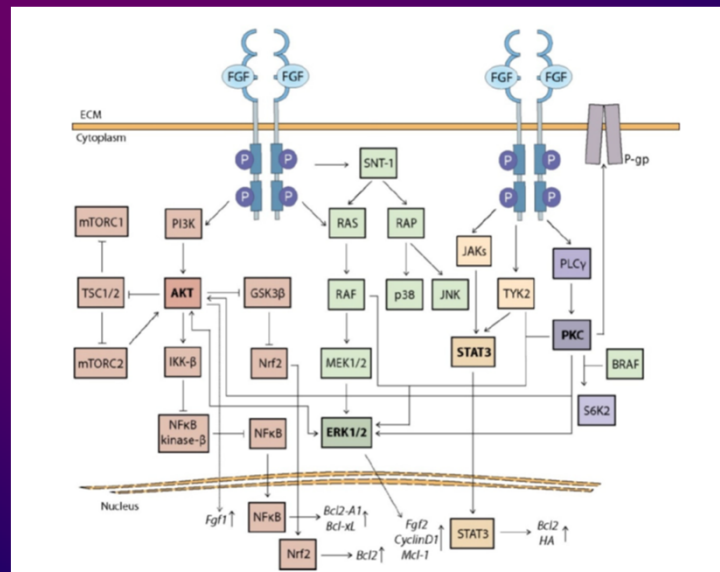
FGFR1/Fgfr1	Fgfr receptor 1	Fgfr	Fgfr-like gene
		FGFR2	Fgfr-like tyrosine kinase 2
		CEK	Chicken embryo kinase 1
		KAL2	Kallman syndrome 2
		K-sam	KATO-III cell-derived stomach cancer amplified gene
FGFR2/Fgfr2	Fgfr Receptor 2	Bek	Bacterial expressed kinase
		CEK3	Chicken embryo kinase 3
		K-Fgf	K-Fgf receptor
FGFR3/Fgfr3	Fgfr Receptor 3	CEK2	Chicken embryo kinase 2
		Ach	Achondroplasia
FGFR4/Fgfr4	Fgfr Receptor 4	TKF	Tyrosine kinase related to Fibroblast Growth Factor receptor
FGFR5/Fgfr5	Fgfr receptor like 1	Fgfr5	Fgfr receptor 5



FGF subfamily	FGF	Cofactor	Receptor specificity
FGF1 subfamily	FGF1		All FGFRs
	FGF2		FGFR 1c, 2c > 2b, 1b, 4Δ
FGF4 subfamily	FGF4		FGFR 1c, 2c > 3c, 4Δ
	FGF5		
	FGF6		
FGF7 subfamily	FGF3	+ Heparin or Heparan sulfate	FGFR 2b > 1b
	FGF7		
	FGF10		
	FGF22		
FGF8 subfamily	FGF8		FGFR 3c > 4Δ > 2c > 1c >> 3b
	FGF17		
	FGF18		
FGF9 subfamily	FGF9		FGFR 3c > 2c > 1c, 3b >> 4Δ
	FGF16		
	FGF20		
FGF15/19 subfamily	FGF15/19	+βKlotho	FGFR 1c, 2c, 3c, 4Δ
	FGF21		FGFR 1c, 3c
	FGF23	+αKlotho	FGFR 1c, 3c, 4

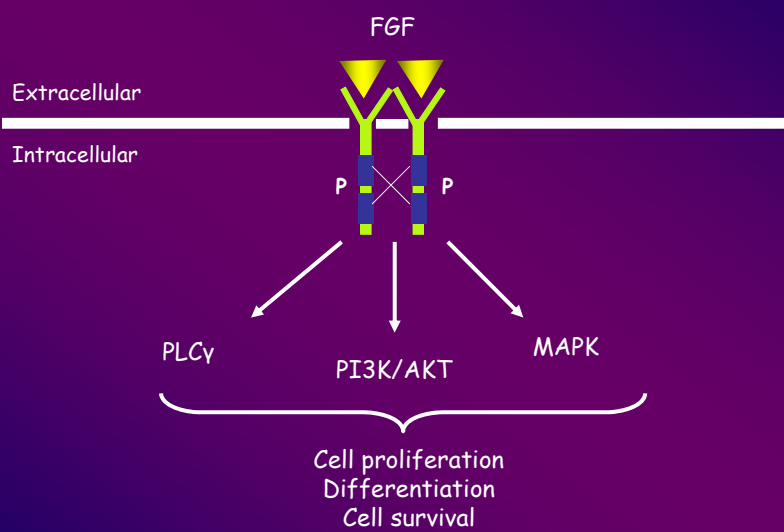
Ornitz and Itoh, Dev. Biol. 2015

### The fibroblast growth factor signaling pathway



Szymczyk et al. 2021 Cancers, 13, 5796

### In addition to the classical receptor tyrosine kinase pathways of FGF...



...there are other FGF pathways: the intracrine pathway.



### The fibroblast growth factors

FGFs	Taille (aa) (MM 18-38 kda)	séquence signal glycosylation	Expression
FGF1/FGFa	155	non	ubiquitaire
FGF2/FGFb	155, 196, 201, 210	non	ubiquitaire
FGF3/int-2	271, 245	oui	embryon, carcinome mammaire
FGF4/hst-1	206	oui	embryon
FGF5	267	oui	embryon, muscle adulte, sarcome de kaposi
FGF6/hst-2	198	oui	muscle squelettique
FGF7/KGF	194	oui	fibroblaste, UVSMC, mésenchyme embryonnaire
FGF8/AIGF	181	oui	ectoderme embryonnaire
FGF9/GAF	174	oui	gliome
FGF10	215	oui	embryon, poumon adulte
FGF11-14/FHF1-4	244, 245, 225, 247	non	CNS (embryon, adulte)
FGF15	218	oui	CNS embryonnaire

Abbreviations: KGF: Keratinocyte growth factor, AIGF: androgen-induced growth factor, GAF: glia activating factor, UVSMC: uterine vascular smooth muscle cells, FHF: fibroblast growth factor homologous factor

#### Activités des FGFs (exogènes)

- facteurs **mitogènes** *in vitro*
- facteurs de **différenciation** et de **transdifférenciation** *in vitro* et *in vivo*
- facteurs de **survie** *in vitro* et *in vivo*
- facteurs **transformant** *in vitro* et **tumorigènes** *in vivo*
- facteurs **angiogéniques**
- facteurs de **réparation** tissulaire

Nuclear  
localization  
sequence

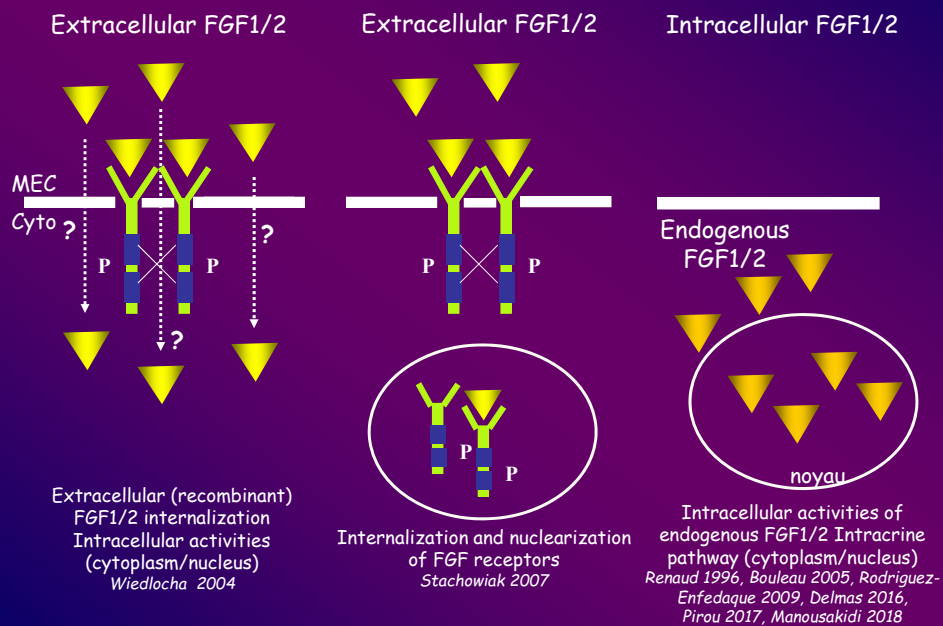
NLS

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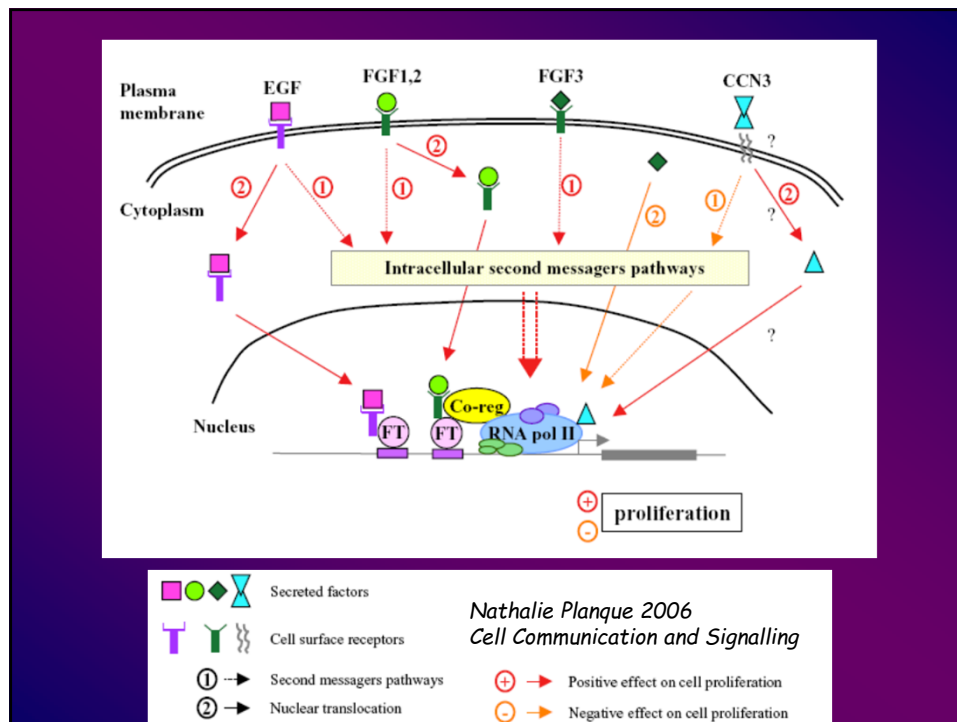
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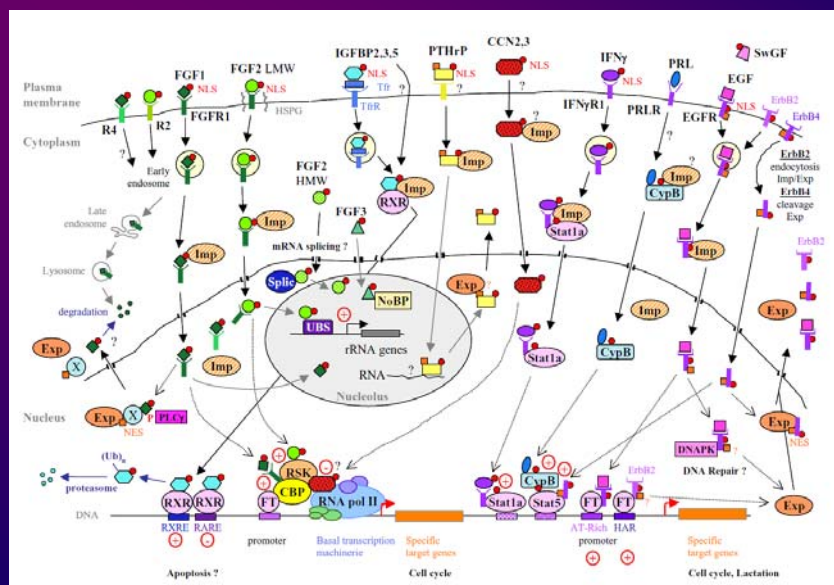
### The Fibroblast Growth Factors 1/2 (FGF1/2)



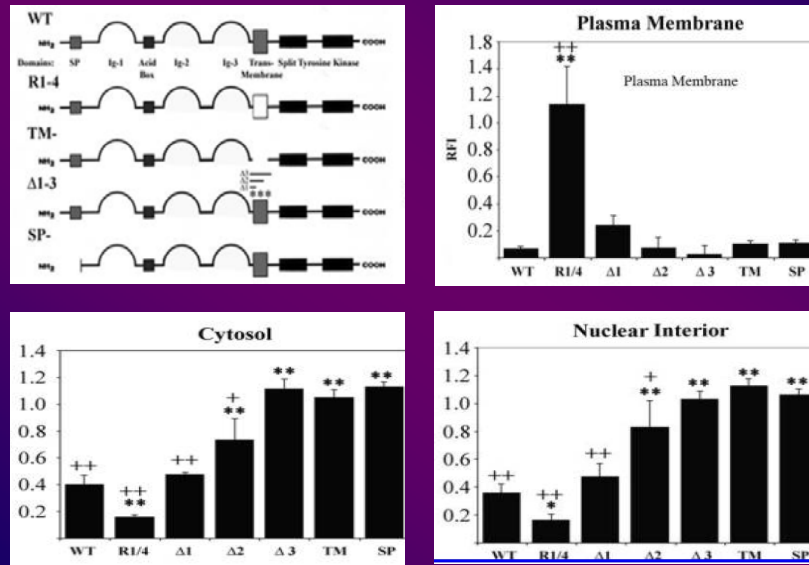




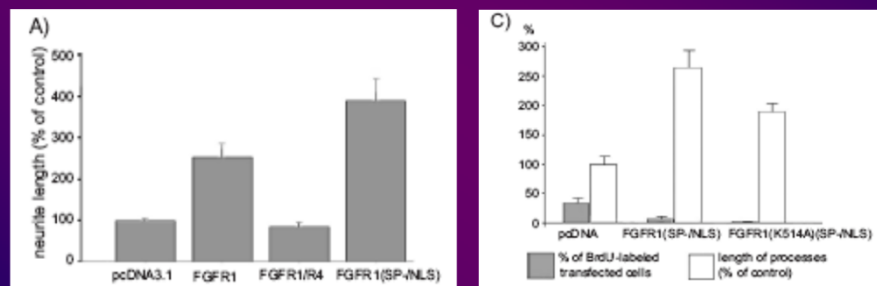
Nathalie Planque 2006 *Cell Communication and Signalling*  
 « Nuclear trafficking of secreted factors and cell-surface receptors: new pathways to regulate cell proliferation and differentiation, and involvement in cancers. »



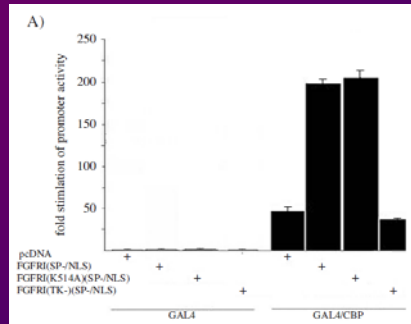
Stachowiak et al. 2007 DNA and Cell Biology  
 « Integrative nuclear signaling in cell development- A role for FGFR1. »



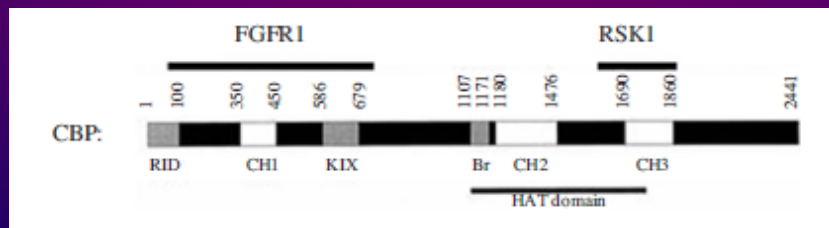
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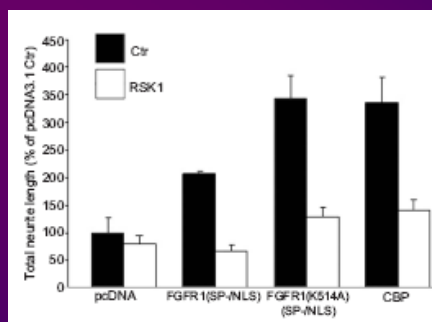
Nuclear FGFR1 stimulates neuronal differentiation of human neuronal primary cells (A) and medulloblastoma TE671 cells (B).  
 The kinase activity of FGFR1 is not required for this activity (K514A mutant TK- of FGFR1 receptor)



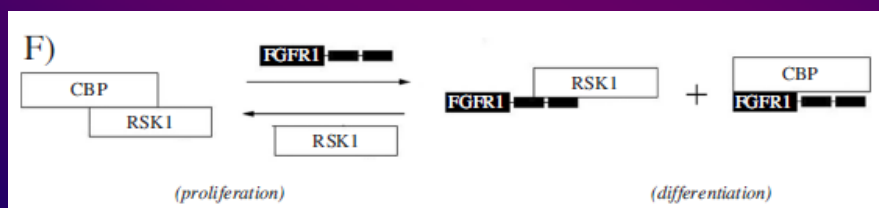
Stachowiak et al. 2007  
DNA and Cell Biology  
« Integrative nuclear signaling in cell development- A role for FGF receptor-1. »



CBP=CREB binding protein: acetyltransferase that activates the transcription of genes which contain AMPc response elements and/or regulates transcription factor activity.  
RSK1=p90<sup>RSK</sup>: serine threonine kinase that regulates several transcription factors.



Stachowiak et al. 2007  
DNA and Cell Biology  
« Integrative nuclear signaling in cell development- A role for FGF receptor-1. »

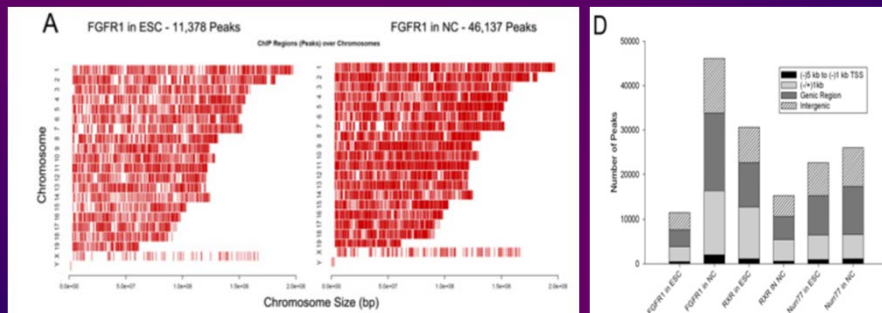


## RESEARCH ARTICLE

# Global Developmental Gene Programming Involves a Nuclear Form of Fibroblast Growth Factor Receptor-1 (FGFR1)

Christopher Terranova<sup>1</sup>, Sridhar T. Narla<sup>1</sup>, Yu-Wei Lee<sup>1</sup>, Jonathan Bard<sup>2</sup>, Abhirath Parikh<sup>3</sup>, Ewa K. Stachowiak<sup>1</sup>, Emmanouel S. Tzanakakis<sup>3</sup>, Michael J. Buck<sup>4</sup>, Barbara Birkaya<sup>3</sup>, Michal K. Stachowiak<sup>1\*</sup>

ESC: pluripotent embryonic stem cell (maintained in presence of LIF)  
NCs: neuronal cells induced by Retinoic Acid (RA)



**Fig 1. Genome-wide analyses of nFGFR1, RXR and Nur77 binding in pluripotent ESCs and RA-induced NCs. (A)** nFGFR1, (B) RXR and (C) Nur77 peaks are present on all chromosomes, in both ESCs and NCs. (D) Genomic distribution of nFGFR1, RXR and Nur77 peaks within proximal promoters (-1 kb to +1 kb relative to TSS), distal promoters (-5 kb to -1 kb relative to TSS), genic and intergenic regions in ESCs and NCs. (E-G) Enrichment of FGFR1, RXR and Nur77 peaks within promoter and genic regions.

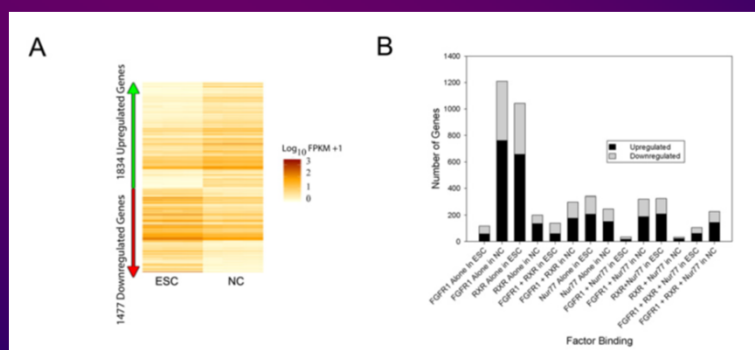
Terranova et al. 2015 PLOSone

## RESEARCH ARTICLE

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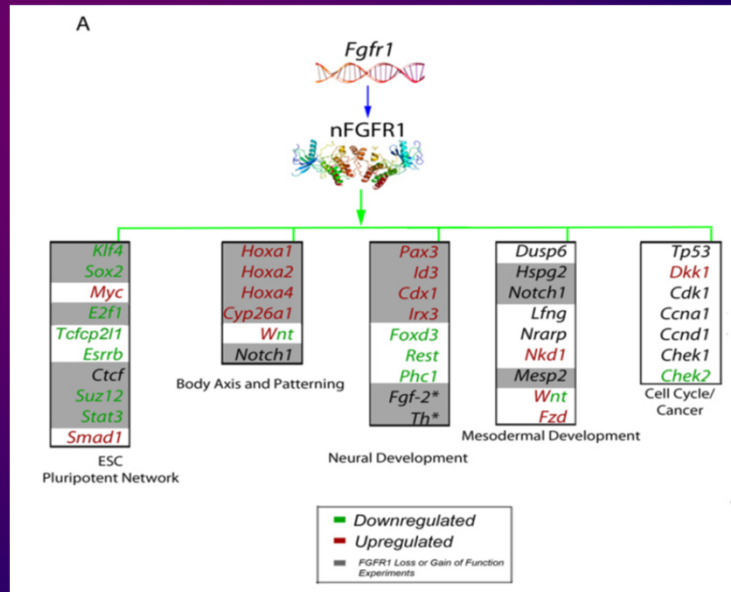
Christopher Terranova<sup>1</sup>, Sridhar T. Narla<sup>1</sup>, Yu-Wei Lee<sup>1</sup>, Jonathan Bard<sup>2</sup>, Abhirath Parikh<sup>3</sup>, Ewa K. Stachowiak<sup>1</sup>, Emmanouel S. Tzanakakis<sup>3</sup>, Michael J. Buck<sup>4</sup>, Barbara Birkaya<sup>3</sup>, Michal K. Stachowiak<sup>1\*</sup>

ESC: pluripotent embryonic stem cell (maintained in presence of LIF)  
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**Fig 3. Binding of nFGFR1, RXR and Nur77 to expressed genes. (A)** Heatmap representation of genes that were differentially expressed in pluripotent ESCs and RA-induced NCs from three independent biological replicates. Out of 14,443 expressed genes, 1,834 were up-regulated and 1,477 were down-regulated in NCs [Fold Change (FC)  $\geq \pm 2.0$  and p-value  $< 0.035$  were considered significant]. Values are displayed as fragments per kb of transcript per million fragments mapped (FPKM). (B) Binding of nFGFR1, RXR and Nur77 within the proximal promoter of differentially regulated genes. In NCs, the population of regulated genes that are targeted by nFGFR1 (2,058 genes) was markedly higher than the population of regulated genes that are not (480 genes).

Terranova et al. 2015 PLOSone



Terranova et al. 2015 PLOSone

### Cancer de la prostate

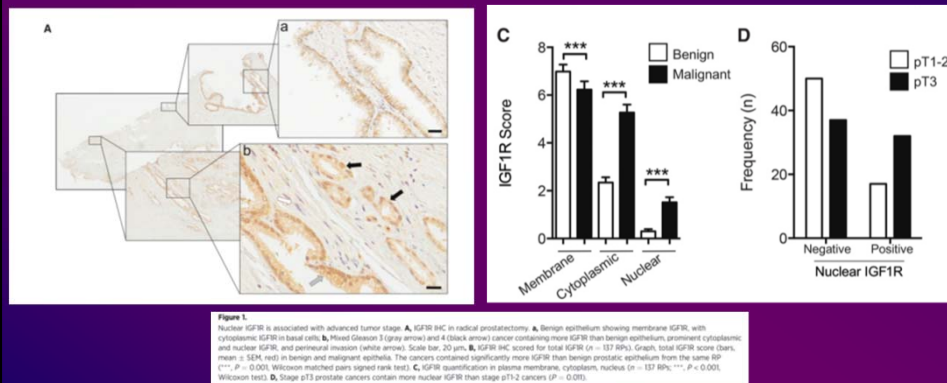
Aleksic et al. 2018

**Molecular Cell Biology**

**Cancer Research**

**Nuclear IGF1R Interacts with Regulatory Regions of Chromatin to Promote RNA Polymerase II Recruitment and Gene Expression Associated with Advanced Tumor Stage**

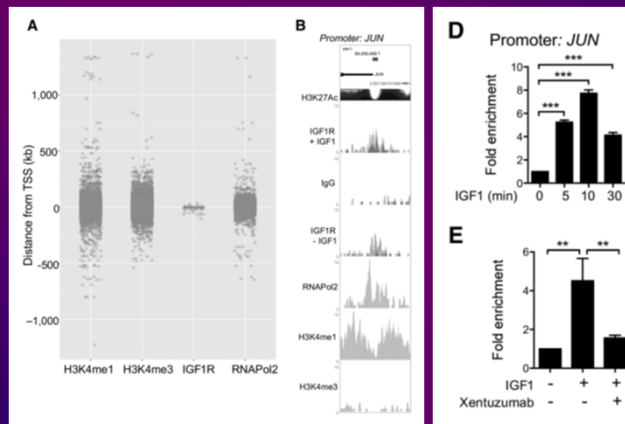
Tamara Aleksic<sup>1</sup>, Nicki Gray<sup>1</sup>, Xiaoning Wu<sup>1</sup>, Guillaume Reunier<sup>1</sup>, Eliot Osher<sup>1</sup>, Jack Mills<sup>1</sup>, Claire Verrier<sup>1</sup>, Richard J. Bryant<sup>1\*</sup>, Cheng Han<sup>1</sup>, Kathryn Hutchinson<sup>1</sup>, Adam G. Lambert<sup>1</sup>, Balraj Kumar<sup>1</sup>, Freddie C. Hamdy<sup>2</sup>, Ulrike Weyer-Camblong<sup>3</sup>, Michael P. Sanderson<sup>4</sup>, Thomas Bogenrieder<sup>5</sup>, Stephen Taylor<sup>2</sup>, and Valentine M. Macaulay<sup>1\*</sup>



**Nuclear IGF1R Interacts with Regulatory Regions of Chromatin to Promote RNA Polymerase II Recruitment and Gene Expression Associated with Advanced Tumor Stage**

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**Figure 2.** IGF1R is recruited to regulatory regions of DNA. **A**, Distance from TSS for ChIP-seq identified peaks. **B**, UCSC browser images. IGF1R-binding sites within JUN and FANCDA/C promoters in DU445 cells treated with or without IGF1 (dark/light gray, duplicate ChIPs), and for IgG, RNAPol2 and H3K4me1/3. H3K27Ac mark, often found near active regulatory elements, from ENCODE (<https://genome.ucsc.edu/ENCODE/>). **C**, ChIP-seq. IGF1R binds to active regulatory elements corresponding to IGF1R-binding peaks in promoters of JUN (lines 1-3) and FANCDA (lines 4-6). White arrow, mobility of free antibody; black, biotinylated probes bound to IGF1R. Mobility is abolished by excess unlabeled probes (lines 3, 5), supporting specificity. No signal in absence of biotinylated probe (line 7). **D**, Serum-starved DU445 cells treated with 50 nmol/L IGF1 for 5-30 minutes, subjected to IGF1R ChIP-qPCR to amplify IGF1R peaks in JUN and FANCDA/C promoters. Graphs, mean  $\pm$  SEM fold enrichment over serum-starved controls. IGF1 increased IGF1R recruitment, peaking at 10 minutes (\*\*\*,  $P < 0.001$ ). **E**, IGF1R ChIP performed as in **D** on serum-starved cells treated with 50 nmol/L IGF1 for 10 minutes alone or with 1 hour 100 nmol/L xentuzumab pretreatment. Graphs, mean  $\pm$  SEM fold enrichment of IGF1R binding to promoters of JUN (left), FANCDA (center), FANCDC (right). \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ .



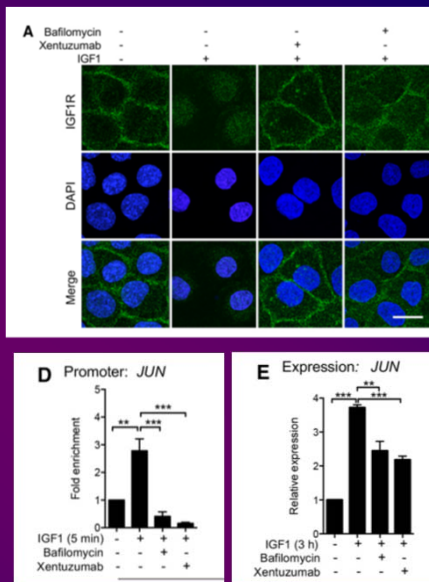
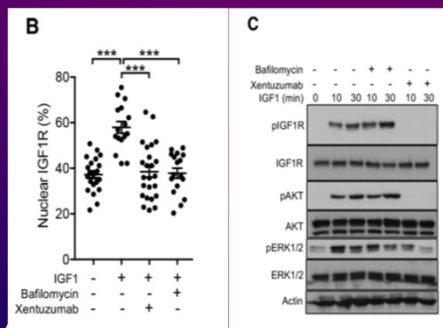
Aleksic et al. 2018

Xentuzumab: Ac monoclonal anti-IGF-1/2

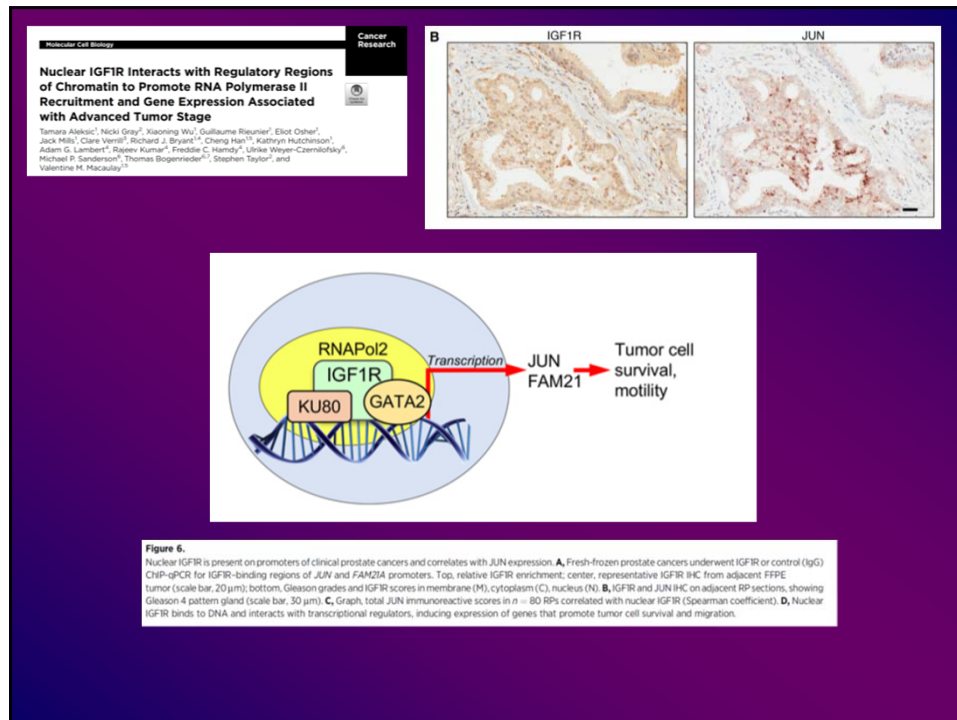
**Nuclear IGF1R Interacts with Regulatory Regions of Chromatin to Promote RNA Polymerase II Recruitment and Gene Expression Associated with Advanced Tumor Stage**

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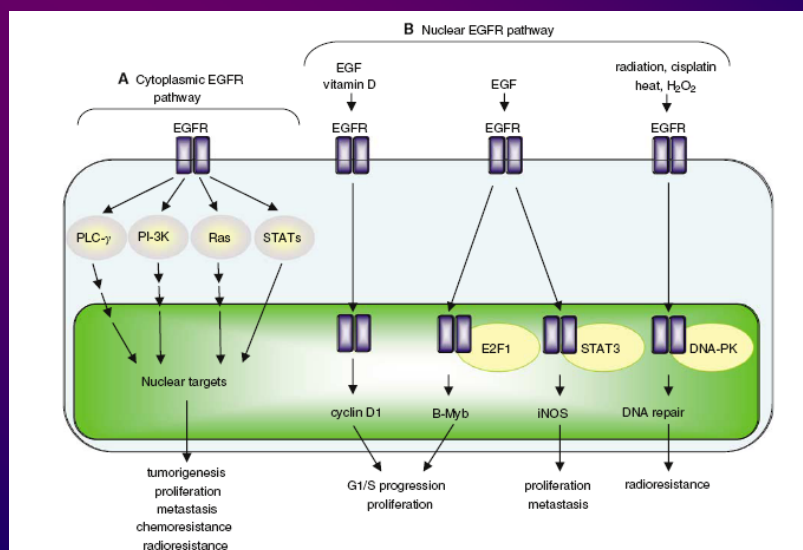
**Figure 4.** IGF1R blockade and inhibition of IGF1R internalization induce comparable suppression of IGF1-induced RNAPol2 recruitment and gene expression. **A-C**, Serum-starved DU445 cells were incubated with 50 nmol/L IGF1 for 30 minutes alone or with 1 hour pretreatment with 100 nmol/L xentuzumab or 50 nmol/L bafilomycin. **A**, Representative IGF1R immunofluorescence images. Scale bar, 20  $\mu$ m. **B**, Quantification of mean  $\pm$  SEM nuclear IGF1R as percentage of total cellular IGF1R (\*\*\*,  $P < 0.001$ ). **C**, Western blot to assess IGF1-induced activation of IGF1R, Akt, and ERK. **D** and **E**, Serum-starved DU445 cells were treated with IGF1 alone or with xentuzumab or bafilomycin. **D**, ChIP-qPCR. IGF1-induced RNAPol2 recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **E**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **F**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **G**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **H**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **I**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **J**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **K**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **L**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **M**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **N**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **O**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **P**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **Q**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **R**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **S**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **T**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **U**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **V**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **W**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **X**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **Y**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin. **Z**, IGF1-induced IGF1R recruitment to JUN and FANCDA/C promoters was attenuated by xentuzumab and bafilomycin.



Xentuzumab: Ac monoclonal anti-IGF-1/2  
Bafilomycin: inhibiteur ATPase vacuolaire, autophagie

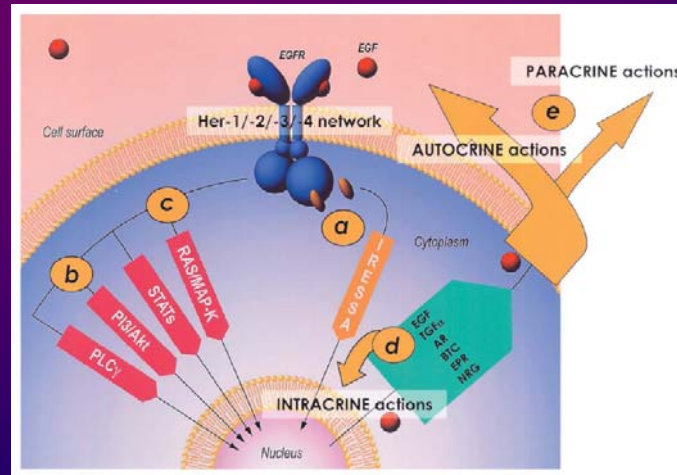


*Lo HW and Hung MC 2006 British Journal of Cancer*  
 « Nuclear EGFR signalling network in cancer: linking EGFR pathway to  
 Cell cycle progression, nitric oxide pathway and patient survival. »





Ferrer-Soler et al. 2007 Int. J. Mol. Med.  
 « An update of the mechanisms of resistance to EGFR-tyrosine kinase inhibitors  
 in breast cancer: Gefitinib (Iressa™)-induced changes in the expression and  
 nucleo-cytoplasmic trafficking of HER-ligands. »



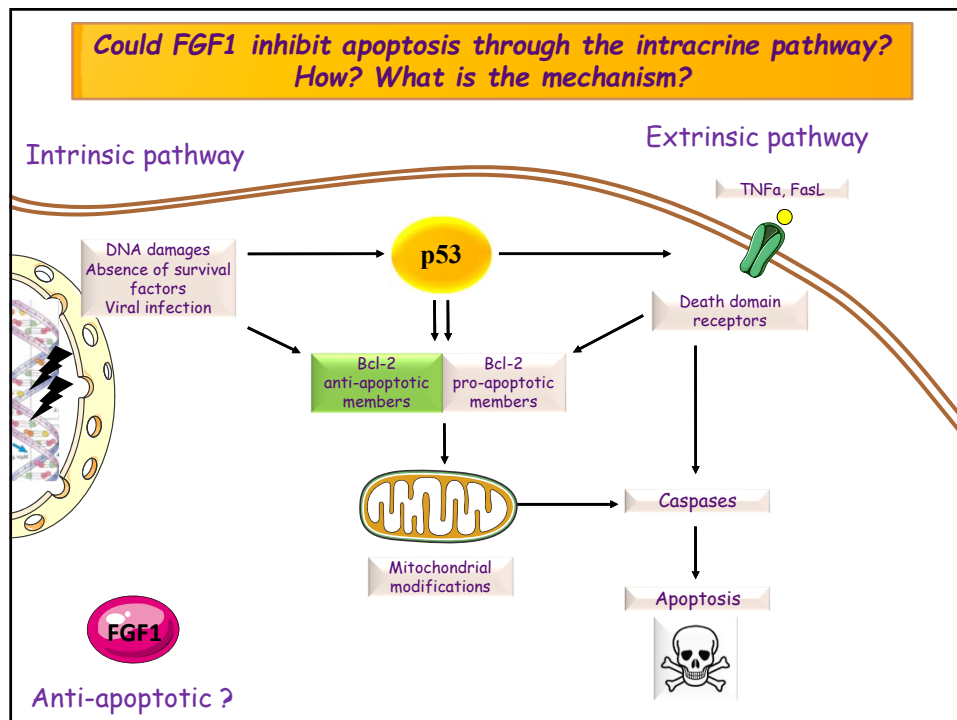
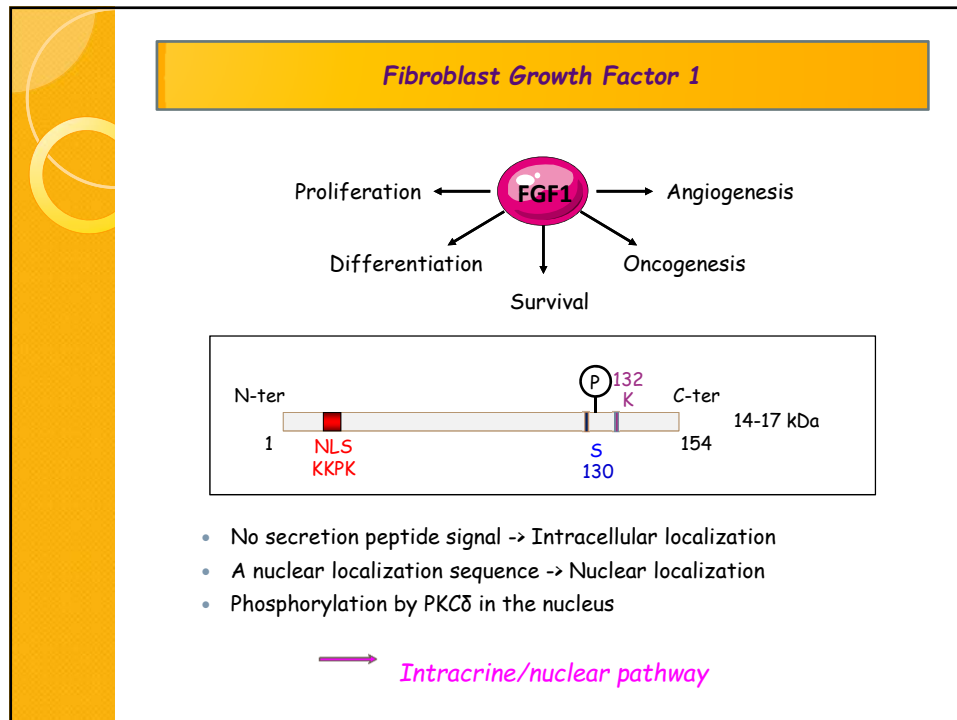
### Study of the FGF1 intracrine pathway

Flore Renaud

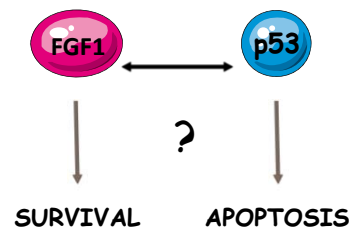
Laboratoire de Génétique et Biologie Cellulaire

LGBC EA4589 UVSQ/EPHE  
Versailles / Montigny le Bretonneux





### Study of the interactions between FGF1 and p53 pathways



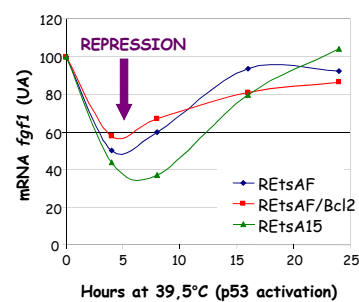
Different cell models:

- Embryonic fibroblasts
- Neuronal cell lines :
  - PC12 (pheochromocytoma),
  - SH-SY5Y (neuroblastoma)
- Ovarian cancer cell lines:
  - COV434 (granulosa),
  - A2780 (serous epithelial)

### p53 represses fgf1 expression in rat embryonic fibroblasts

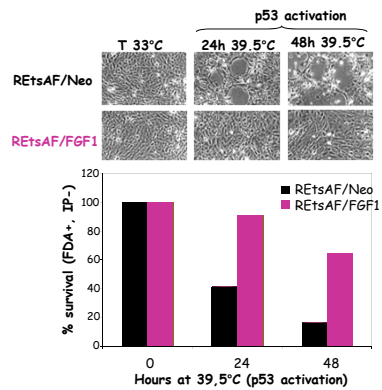
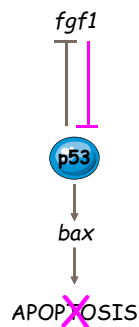


1/ *fgf1* is a p53 target gene.



Bouleau et al. *Oncogene* 2005

**FGF1 inhibits p53 pro-apoptotic and anti-proliferative activities by an intracrine pathway in rat embryonic fibroblasts**

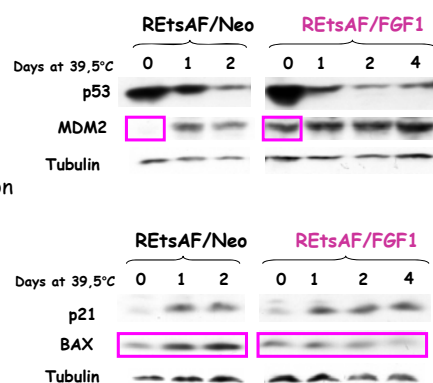
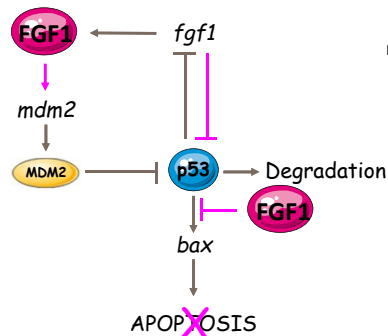


1/ *fgf1* is a p53 target gene.

2/ FGF1 inhibits p53 pro-apoptotic and anti-proliferative activities by an intracrine pathway.

Bouleau et al. *Oncogene* 2005

**FGF1 inhibits p53 pro-apoptotic and anti-proliferative activities by an intracrine pathway in rat embryonic fibroblasts**



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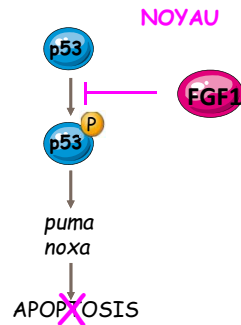
2/ FGF1 inhibits p53 pro-apoptotic and anti-proliferative activities by an intracrine pathway.

- ✓ FGF1 increases *mdm2* expression and p53 degradation
- ✓ FGF1 inhibits some p53 transcriptional activities (*bax*)

Bouleau et al. *Oncogene* 2005

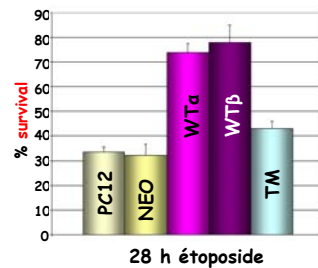
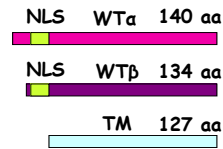
### Intracellular FGF1 inhibits p53-dependent apoptosis in PC12 cells by an intracrine/nuclear pathway

1/ Extracellular and intracellular FGF1 inhibits p53-dependent apoptosis.



2/ FGF1 nuclearization is required for its neurotrophic and anti-apoptotic activities.

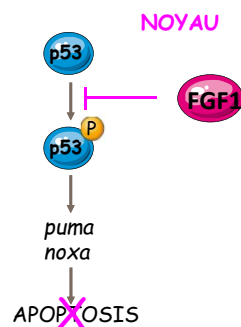
Different forms of FGF1



Bouleau et al. 2007  
Rodriguez-Enfedaque et al. 2009

### Intracellular FGF1 inhibits p53-dependent apoptosis in PC12 cells by an intracrine/nuclear pathway

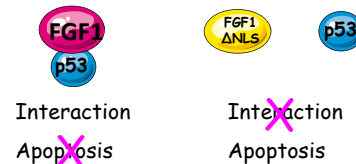
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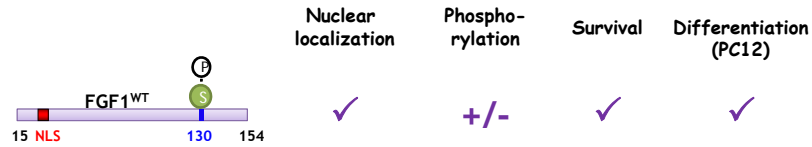
3/ FGF1 interacts with p53 (directly or indirectly?).

The nuclear localization sequence is required for this interaction.



Bouleau et al. 2007  
Rodriguez-Enfedaque et al. 2009

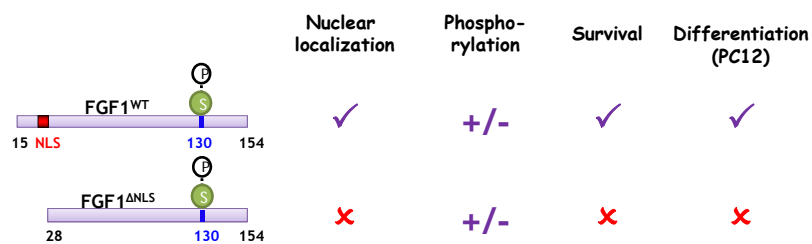
**Both nuclearization and phosphorylation regulate FGF1 activities in neuronal cell lines (PC12, SH-SY5Y)**



In both neuronal cell lines (PC12 and SH-SY5Y cell lines):  
1/ Intracellular FGF1 inhibits p53-dependent apoptosis

*Delmas et al. 2016, Pirou et al. 2017*

**Both nuclearization and phosphorylation regulate FGF1 activities in neuronal cell lines (PC12, SH-SY5Y)**



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1/ Intracellular FGF1 inhibits p53-dependent apoptosis  
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*Delmas et al. 2016, Pirou et al. 2017*

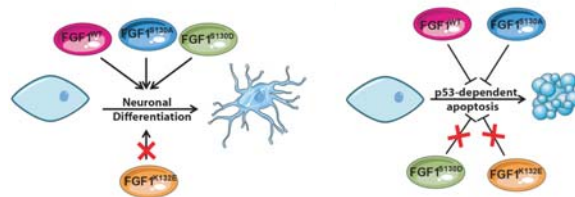
**Both nuclearization and phosphorylation regulate FGF1 activities in neuronal cell lines (PC12, SH-SY5Y)**

	Nuclear localization	Phosphorylation	Survival	Differentiation (PC12)
 FGF1 <sup>WT</sup> 15 NLS 130 154	✓	+/-	✓	✓
 FGF1 <sup>ΔNLS</sup> 28 130 154	✗	+/-	✗	✗
 FGF1 <sup>S130A</sup> 15 NLS 130 154	✓	Inhibits	✓	✓
 FGF1 <sup>S130D</sup> 15 NLS 130 154	✓	Mimics	✗	✓

In both neuronal cell lines (PC12 and SH-SY5Y cell lines):  
 1/ Intracellular FGF1 inhibits p53-dependent apoptosis  
 2/ FGF1 nuclearization is required for its anti-apoptotic activity.  
 3/ FGF1 phosphorylation inhibits its anti-apoptotic activity

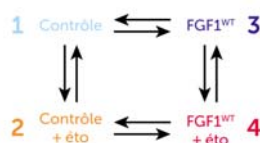
Delmas et al. 2016, Pirou et al. 2017

**Study of the nuclear FGF1 pathway:  
Transcriptome analysis of PC12 cells +/- FGF1 +/- p53**



RNA-seq analysis in PC12 cells

P < 0,01	FGF1 <sup>WT</sup> VS Ctrl	Ctrl + étio VS Ctrl	FGF1 <sup>WT</sup> + étio VS FGF1 <sup>WT</sup>	FGF1 <sup>WT</sup> + étio VS Ctrl + étio
Up-regulated genes number	1709	1891	1065	284
Down-regulated genes number	1780	2591	1381	340



**Genes and/or signaling pathway regulated:**

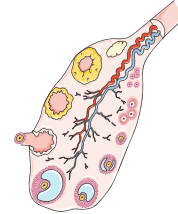
- *fgf1* (amplification loop)
- Transcription factors (*Etv1*, *Egr1*, *c-jun*, ...)
- Wnt/ $\beta$ -catenin pathway (*APC*, *DKK3*, *Wnt7a*)
- Jak/Stat pathway.

Caroline Pirou, PhD 2016

### Study of FGF1 activity in chemoresistance of ovarian cancers

#### In ovarian cancer:

- *fgf1* overexpression correlated with high grade and chemoresistance of ovarian tumor cells (Smith et al. 2012).
- FGF1 increases tumor growth *in vivo* (King et al. 2014).



#### Study of FGF1 activity (and mechanism) in the resistance of ovarian tumor cells to etoposide or cisplatin treatment

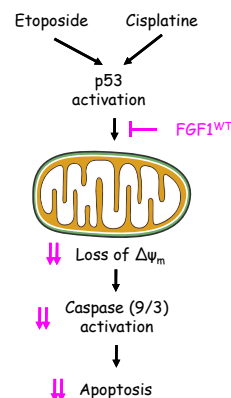
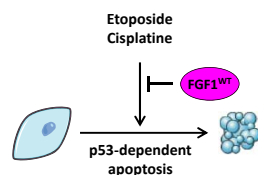
##### 2 ovarian tumor cell lines:

- COV434: cell line issued from a granulosa tumor
- A2780: cell line issued from a serous epithelial tumor (used by Smith and King teams)

### Study of FGF1 activity in chemoresistance of ovarian cancers

#### In COV434 cells:

- Extracellular and intracellular FGF1 protects COV434 cells from etoposide- and cisplatin-induced apoptosis.

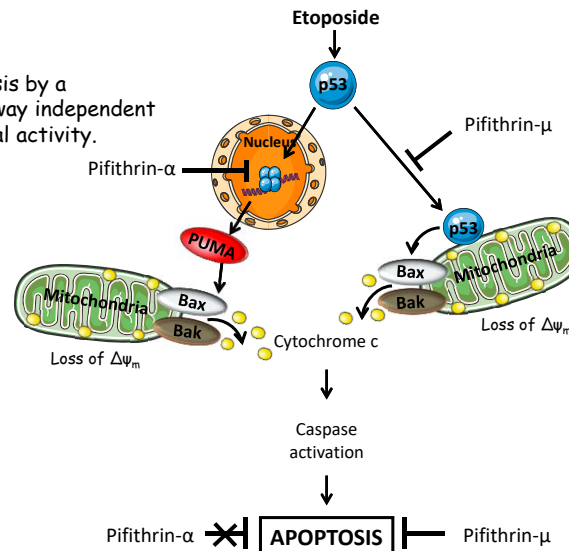


Manousakidi et al. 2018

### Study of FGF1 activity in chemoresistance of ovarian cancers

#### In COV434 cells:

- p53 induces apoptosis by a mitochondrial pathway independent of its transcriptional activity.

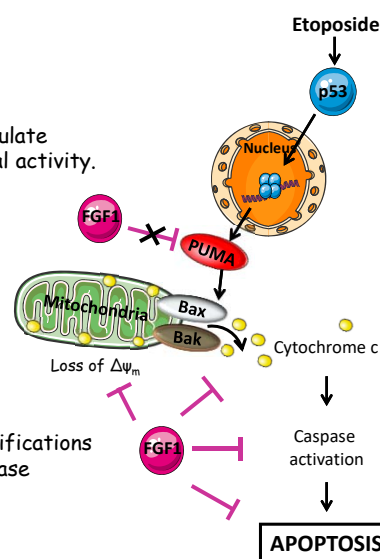


Manousakidi et al. 2018

### Study of FGF1 activity in chemoresistance of ovarian cancers

#### In COV434 cells:

- FGF1 does not regulate p53 transcriptional activity.



- FGF1 inhibits:
  - mitochondrial modifications
  - cytochrome c release
  - caspase activation

Manousakidi et al. 2018

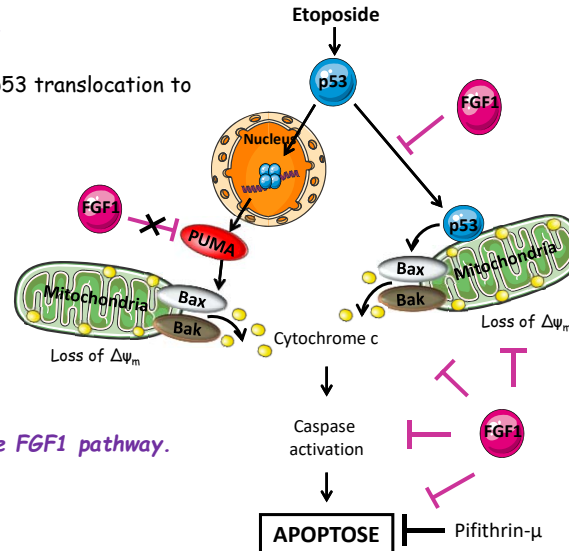


### Study of FGF1 activity in chemoresistance of ovarian cancers

In COV434 cells:

- FGF1 inhibits p53 translocation to mitochondria.

→ New intracrine FGF1 pathway.



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LGBC  
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et biologie cellulaire  
UNIVERSITÉ DE  
VERSAILLES  
ST-QUENTIN-EN-YVELINES

« Etude des régulations de la voie mitochondriale de l'apoptose  
par la voie d'action intracrine du FGF1 »

Groupe « Biologie cellulaire de l'apoptose » LGBC EA4589 UVSQ/EPHE


Flore Renaud

Sylvina Bouleau  
Aida Rodriguez-Enfedaque  
Hélène Grimal


Jean-Luc Vayssière  
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Caroline Pirou  
Nadège Jah  
Fatemeh Torbati


Nathalie Leleu  
Sévasti Manousakidi  
Arnaud Guillaume




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***From the study of the intracrine pathway of FGF1  
to the study of kidney cancer predisposition genes***


Flore Renaud, PhD, HDR, MCF EPHE  
[flore.renaud-paitra@ephe.psl.eu](mailto:flore.renaud-paitra@ephe.psl.eu); [flore.renaud-paitra@gustaveroussy.fr](mailto:flore.renaud-paitra@gustaveroussy.fr)

Laboratoire de Génétique et Biologie Cellulaire, LGBC EA4589 UVSQ/EPHE  
Versailles/ Montigny le Bretonneux (1997-2017)


UMR 1186 INSERM : « Immunologie intégrative des tumeurs et génétique oncologique » (2017-2019)  
 UMR 9019 CNRS : “Genome Integrity and Cancers” (since 2020),  
 Team “Mechanisms of DNA Repair and Carcinogenesis” (Murat Saparbaev)  
 Kidney cancer research group (Stéphane Richard, Sophie Gad)  
 Gustave Roussy Institut, Villejuif




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FONDATION ARC  
Pour la RECHERCHE  
sur le CANCER



PREDIR



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LA LIGUE  
CONTRE LE CANCER