

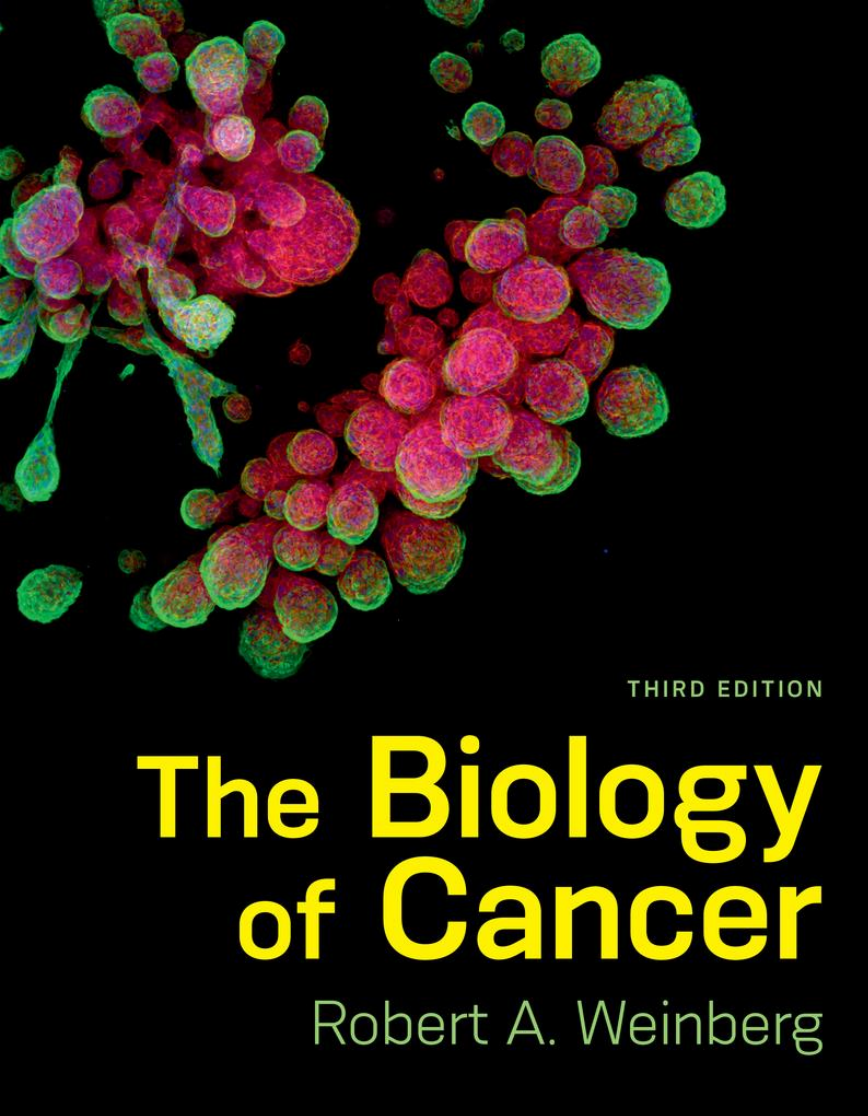


Chapter 07

TUMOR SUPPRESSOR GENES

Chapter 08

RB AND CONTROL OF THE CELL CYCLE CLOCK



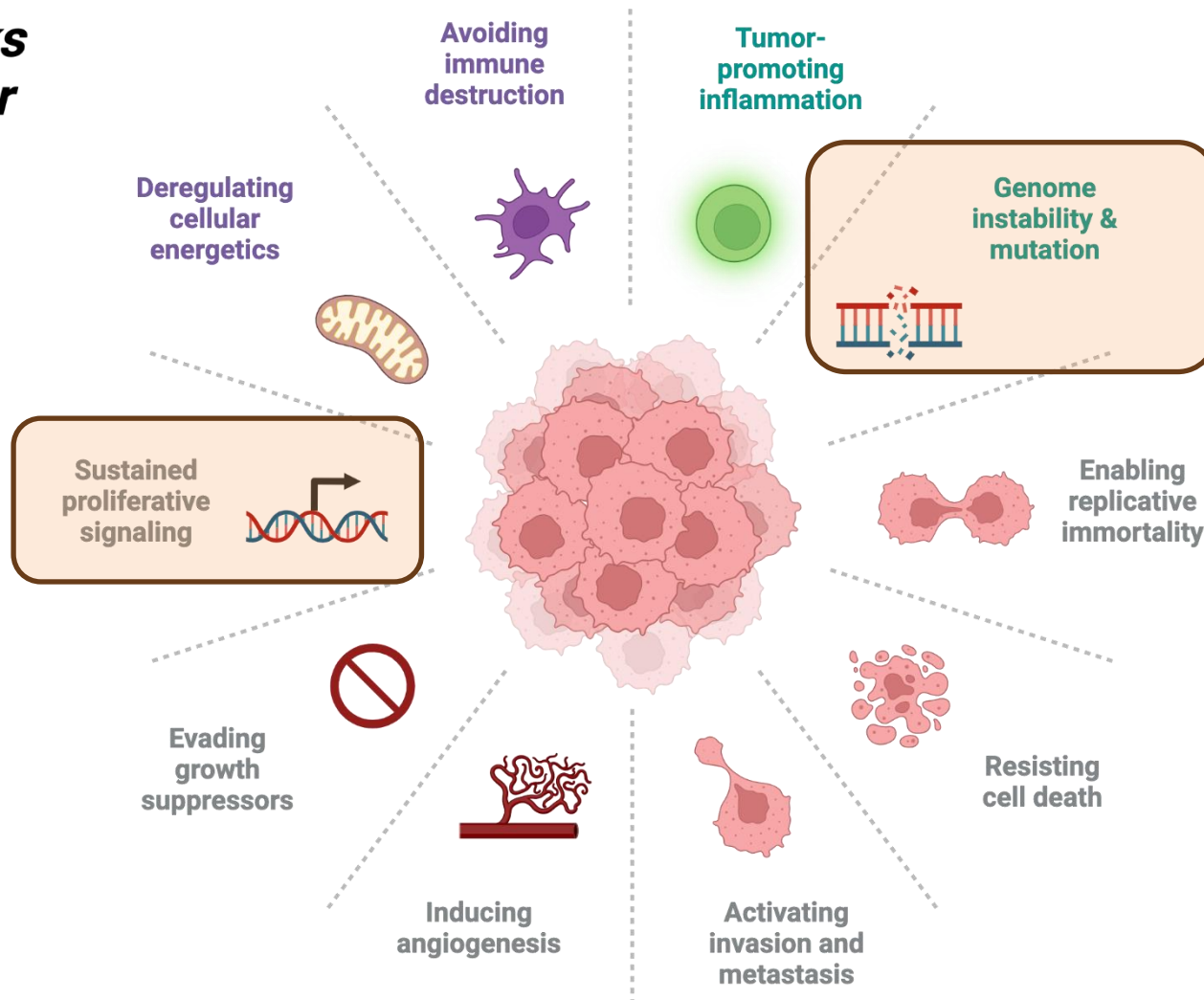
Hee Won Yang (hy2602@cumc.columbia.edu)

Organization of Lecture

- n Proto-oncogenes vs. Tumor suppressor genes
- n The retinoblastoma protein
- n The p53 gene
- n Targeting the Rb pathway in cancer

Hallmarks of cancer

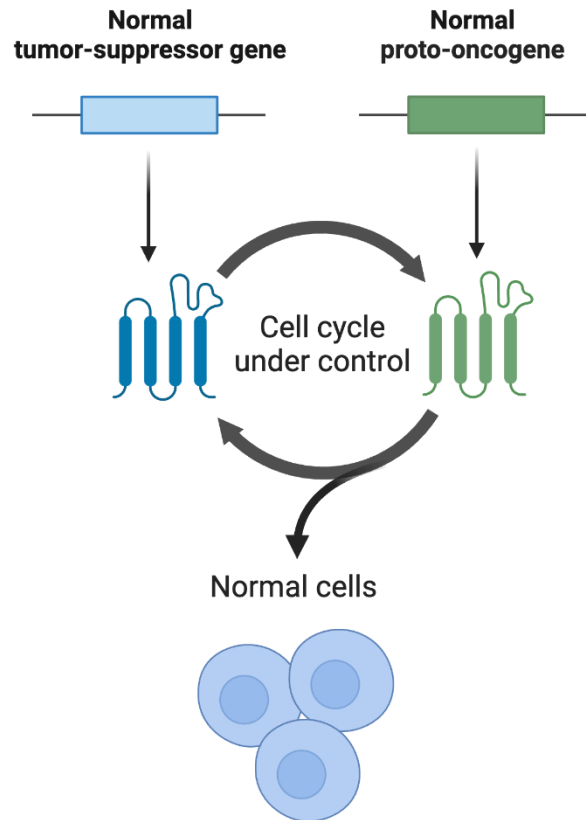
Hallmarks of Cancer



Original hallmarks
Enabling factors
Emerging hallmarks

Proto-oncogenes vs. Tumor suppressor genes

Normal Cell Division



Proto-oncogenes vs. Tumor suppressor genes

- Some are altered in a restricted set of tumor types
 - e.g., the APC (adenomatous polyposis coli) tumor suppressor in colorectal carcinoma
- Others are altered in a broad spectrum of tumor types
 - e.g., p53 tumor suppressor and the Ras proto-oncogenes
- The importance of tumor gene “pathways”
 - the Rb and p53 pathways

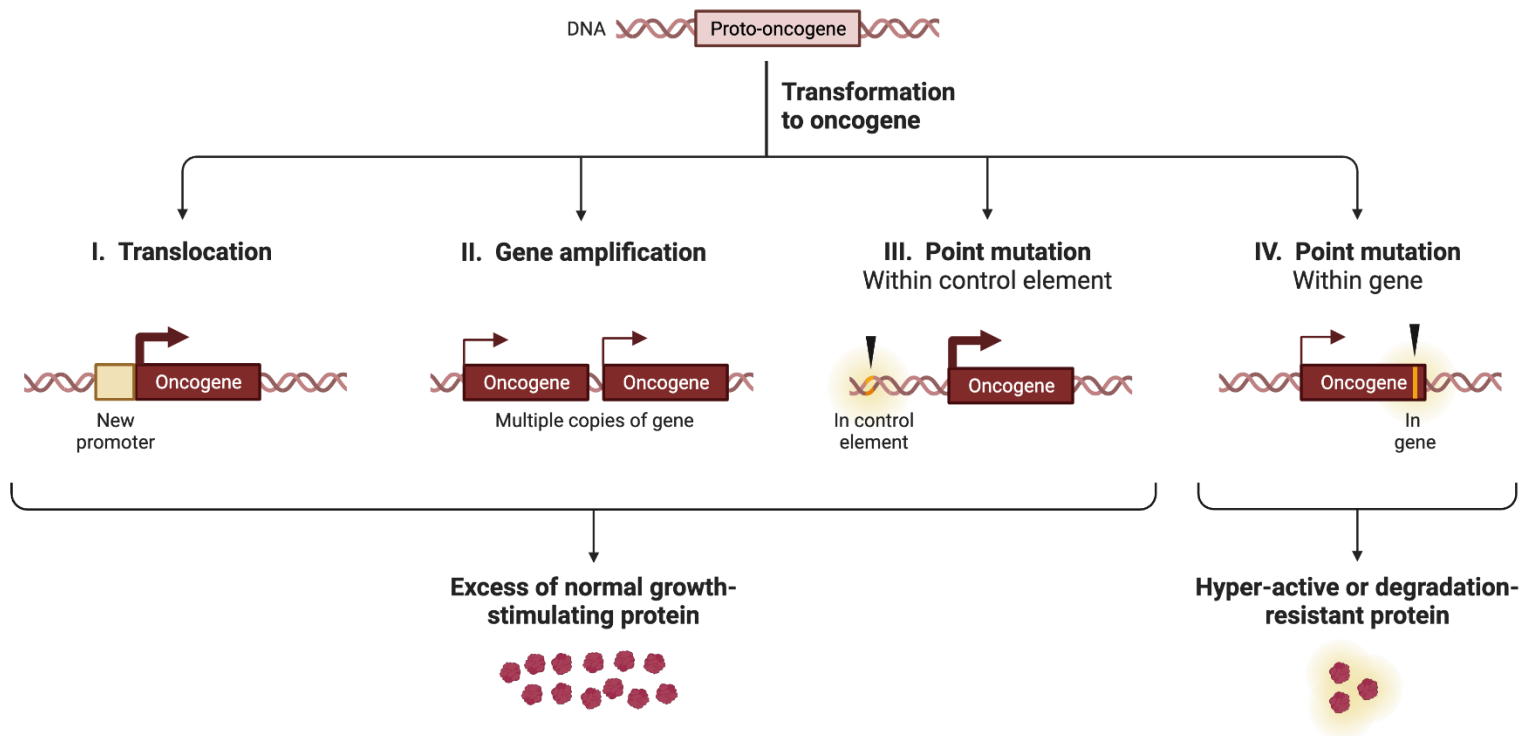
Proto-oncogenes vs. Tumor suppressor genes

A **proto-oncogene** promotes cancer when its function is malignantly activated

- An activated proto-oncogene contributes to tumorigenesis by "**gain-of-function**"
- Thus, an activated proto-oncogene is genetically dominant at the cellular level
 - an activated oncogene can elicit a new phenotype (tumorigenesis) even in the presence of the corresponding wild type allele

Proto-oncogenes vs. Tumor suppressor genes



Methods of Oncogene Activation in Cancer



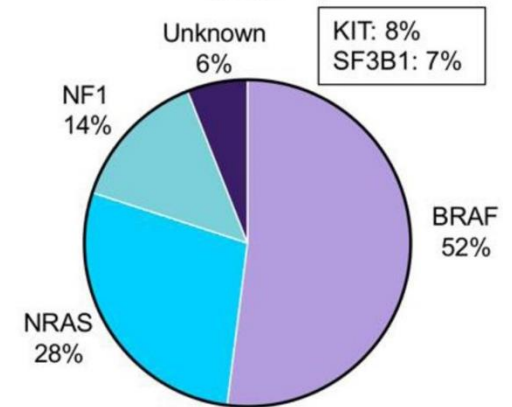
- gene modification, leading to overexpression (e.g., protein overexpression), leading to gene dysregulation or overexpression
 - Ras gene amplification in many cancer types
 - c-Myc gene translocation in Burkitt's lymphoma

Proto-oncogene mutations in Melanoma

Estimated New Cases

			Males	Females			
Prostate	288,300	29%			Breast	297,790	31%
Lung & bronchus	117,550	12%			Lung & bronchus	120,790	13%
Colon & rectum	81,860	8%			Colon & rectum	71,160	8%
Urinary bladder	62,420	6%			Uterine corpus	66,200	7%
Melanoma of the skin	58,120	6%			Melanoma of the skin	39,490	4%
Kidney & renal pelvis	52,360	5%			Non-Hodgkin lymphoma	35,670	4%
Non-Hodgkin lymphoma	44,880	4%			Thyroid	31,180	3%
Oral cavity & pharynx	39,290	4%			Pancreas	30,920	3%
Leukemia	35,670	4%			Kidney & renal pelvis	29,440	3%
Pancreas	33,130	3%			Leukemia	23,940	3%
All Sites	1,010,310	100%			All Sites	948,000	100%

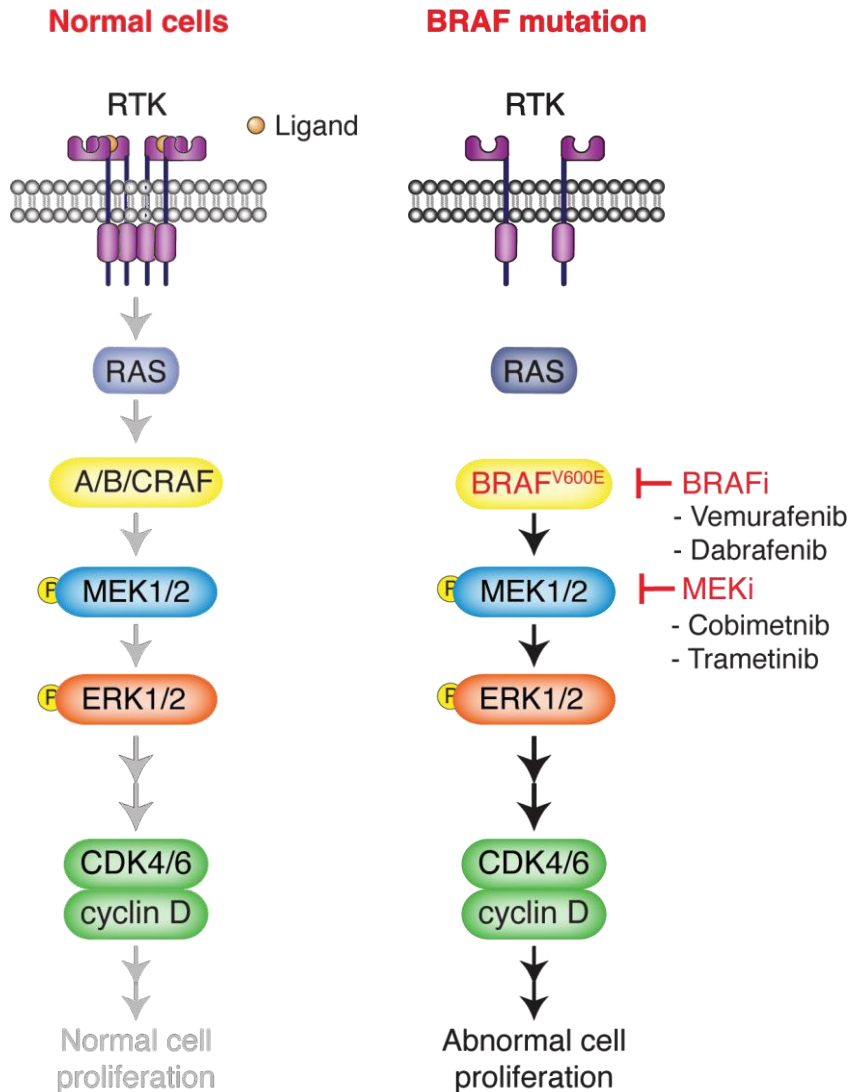
Cutaneous Melanoma: TCGA n=331



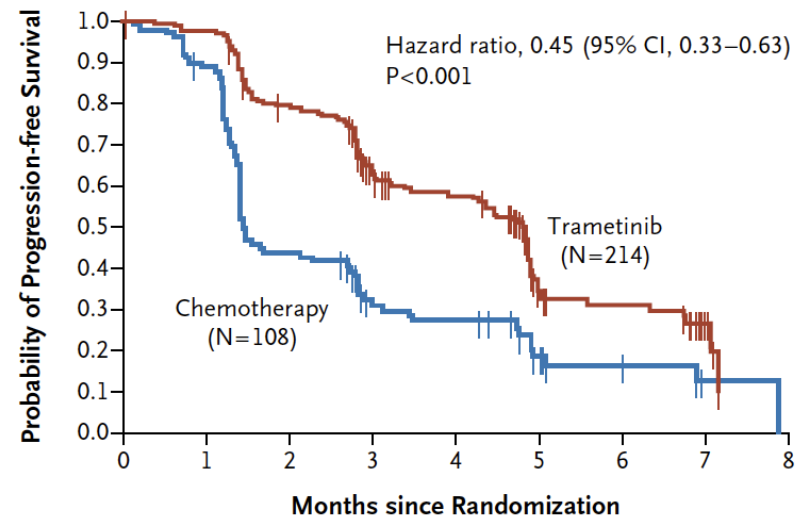
Cancer statistics, 2023

Nassar et al., 2020

BRAF V600 mutation in melanoma cells

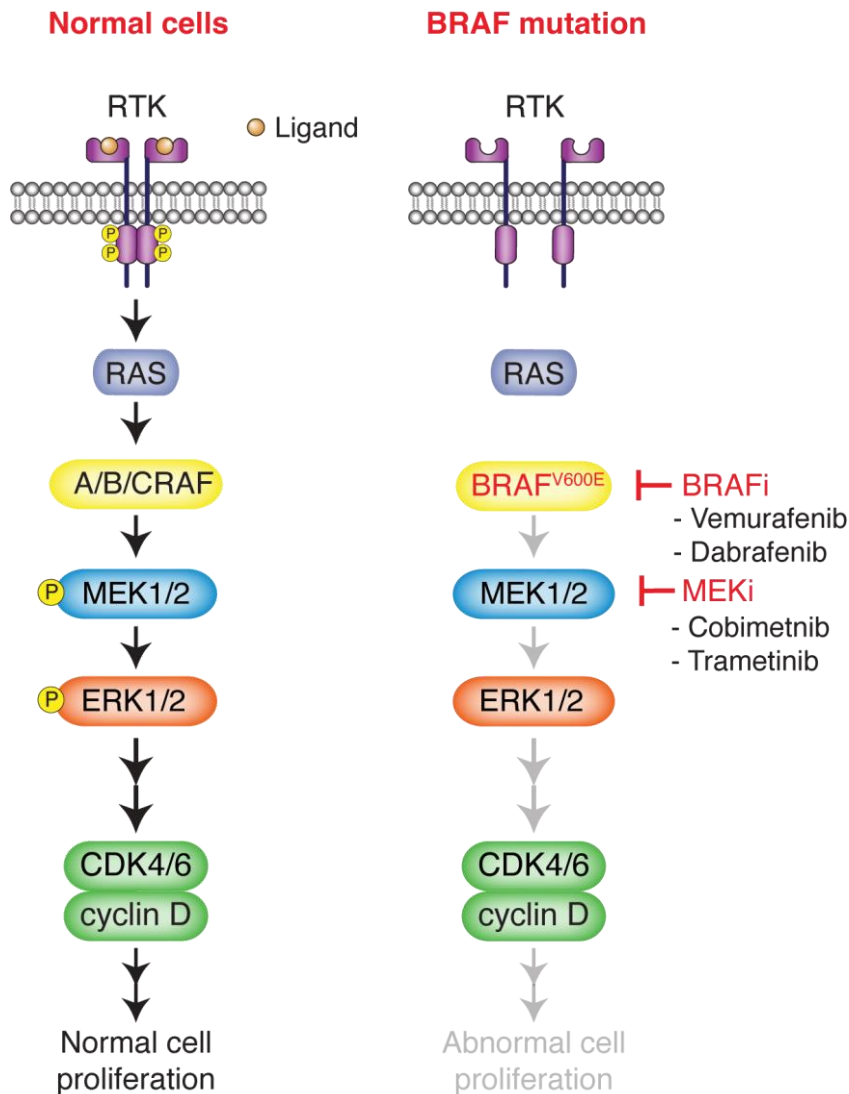


Progression-free Survival



Flaherty et al., 2012

BRAF V600 mutation in melanoma cells



Treated with BRAF inhibitor
(Response rate >50%; Response duration: ~6 months)

38-year-old man with BRAF-mutant melanoma

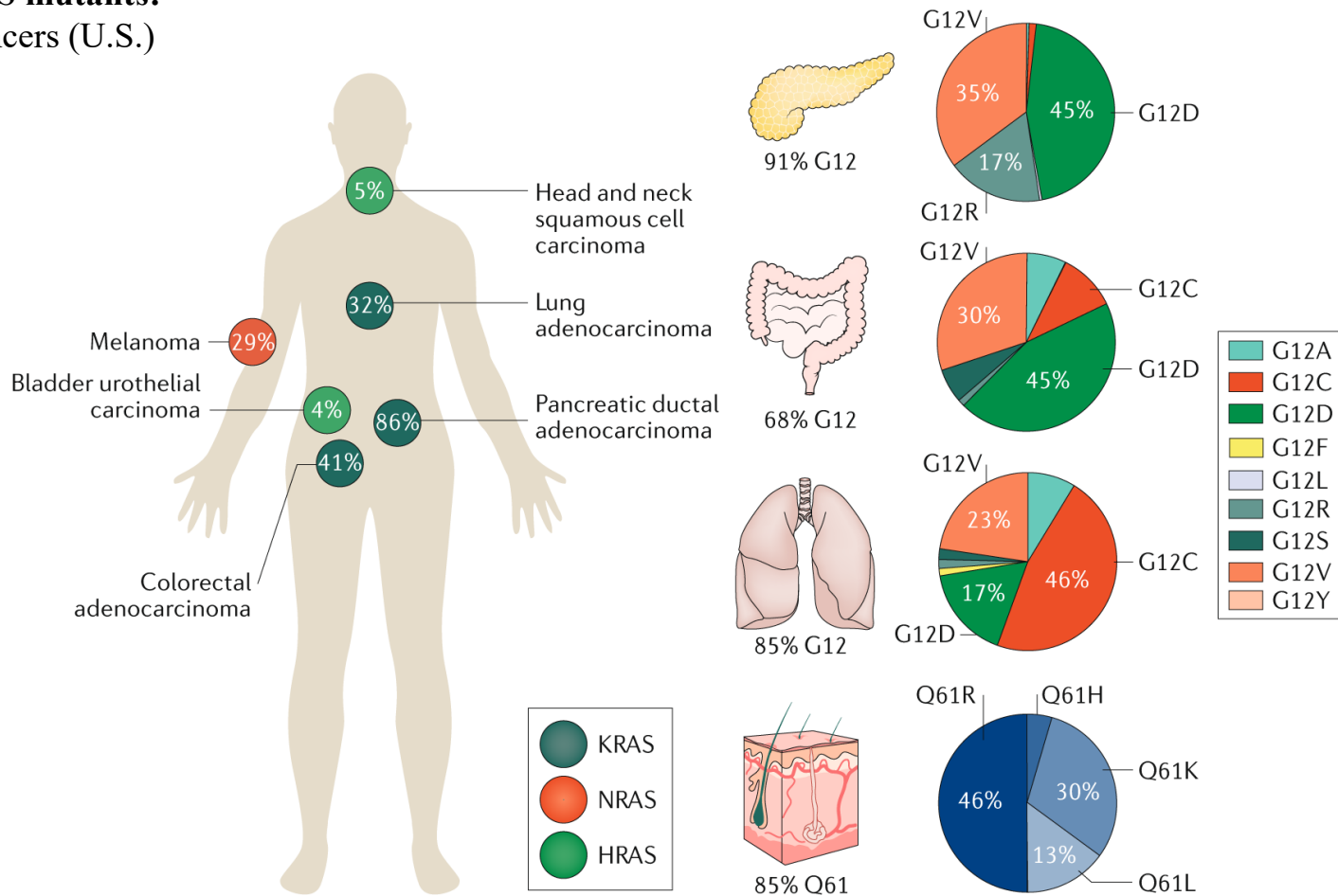
Wagle et al., 2011

About 40% of patients develop resistance to BRAF/MEK inhibitor within a year.

RAS is one of the most frequently mutated genes in human cancers

Diverse RAS mutants:

~30% of cancers (U.S.)



Proto-oncogenes vs. Tumor suppressor genes

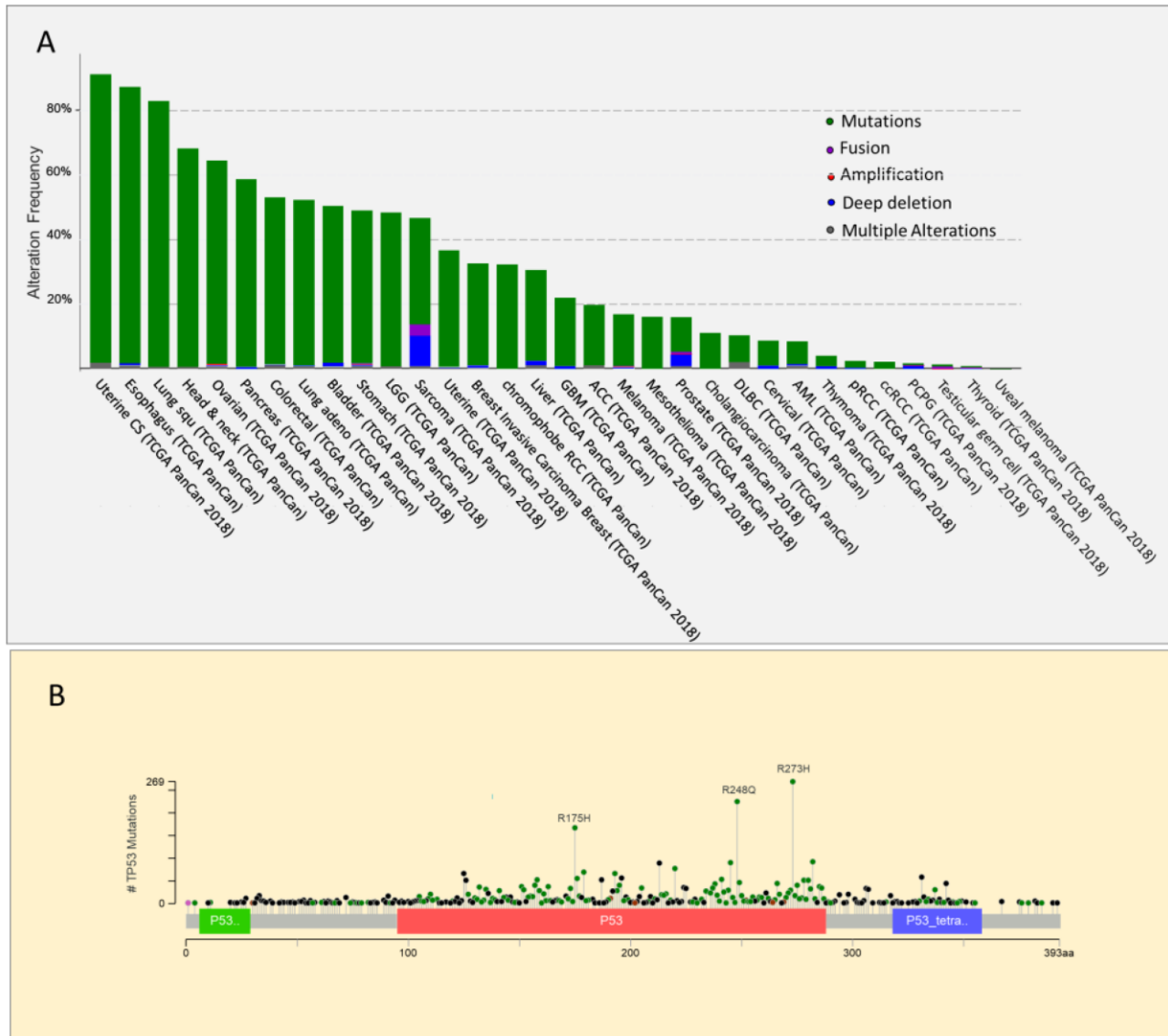
A tumor suppressor gene promotes cancer when its function is malignantly inactivated

- A tumor suppressor contributes to tumorigenesis by "loss-of-function"
- In most instances, an inactivated tumor suppressor gene is genetically recessive at the cellular level.
 - It will not promote tumorigenesis in diploid cells unless the other (wildtype) allele is also lost or inactivated
 - Some exceptions:
 - dominant-negative p53 mutations
 - "haploinsufficient" tumor suppressor genes

Proto-oncogenes vs. Tumor suppressor genes

- In this lecture we will focus on...
 - the retinoblastoma susceptibility (Rb) gene
 - the p53 tumor suppressor gene
- Genetic properties
- Biochemical functions of their protein products
- the p53 and Rb tumor suppressor “pathways”

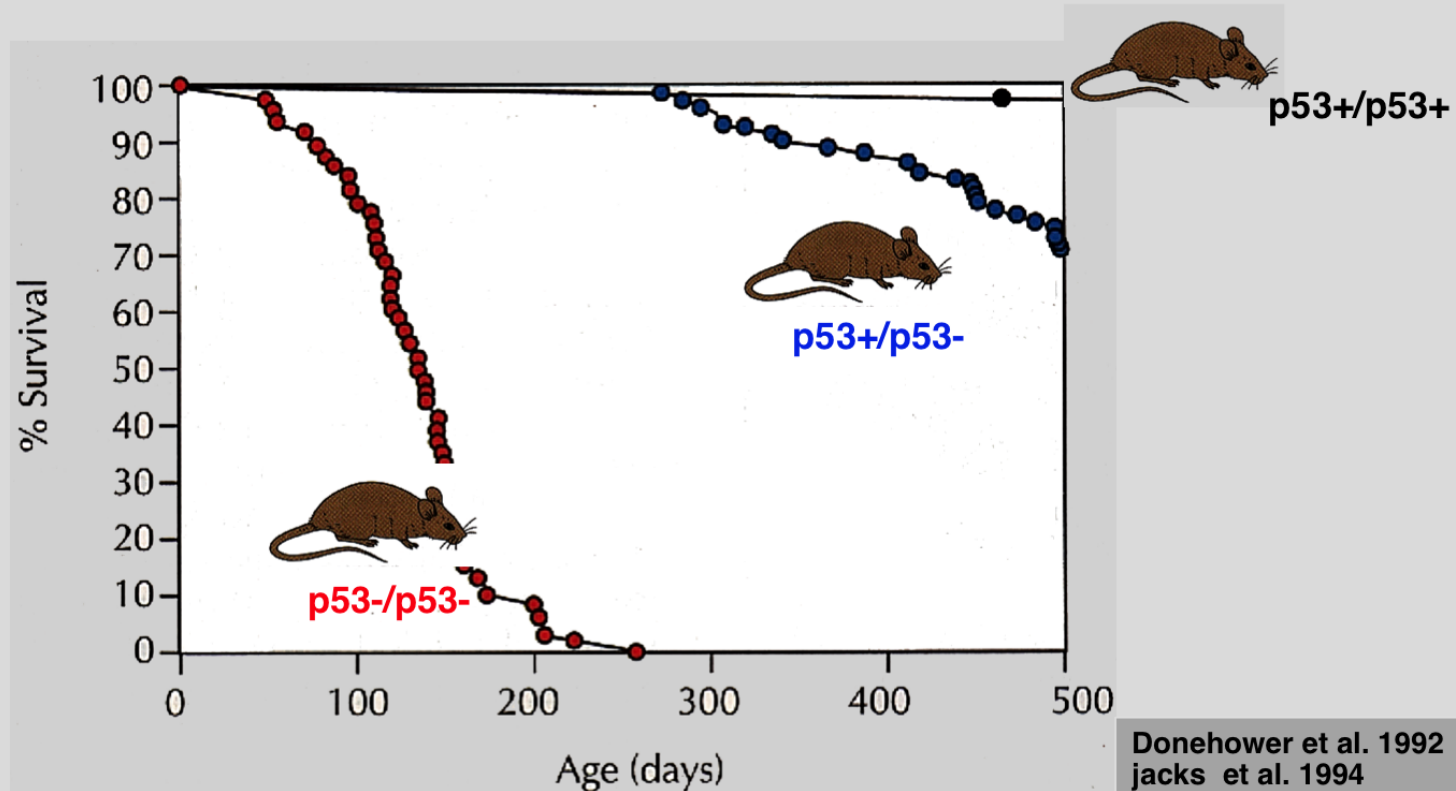
p53 is the most frequently mutated gene in cancer



The p53 gene

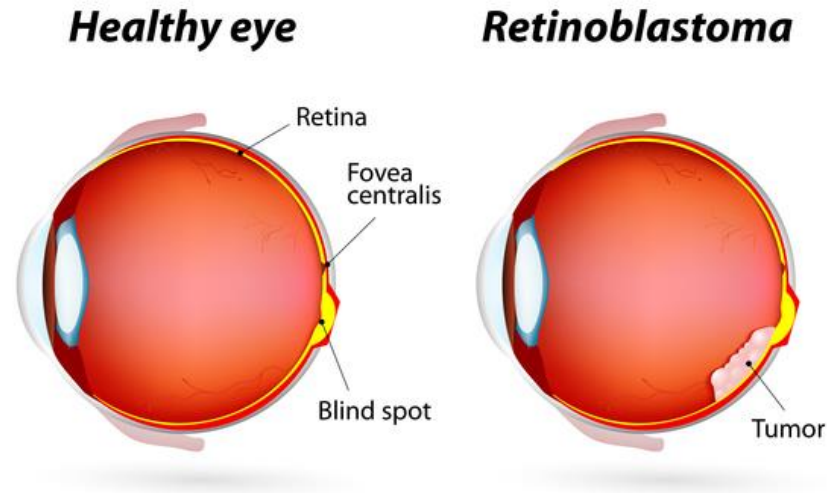
p53: A TUMOR SUPPRESSOR GENE ? (VI)

Mice deficient for p53 are prone to develop cancer



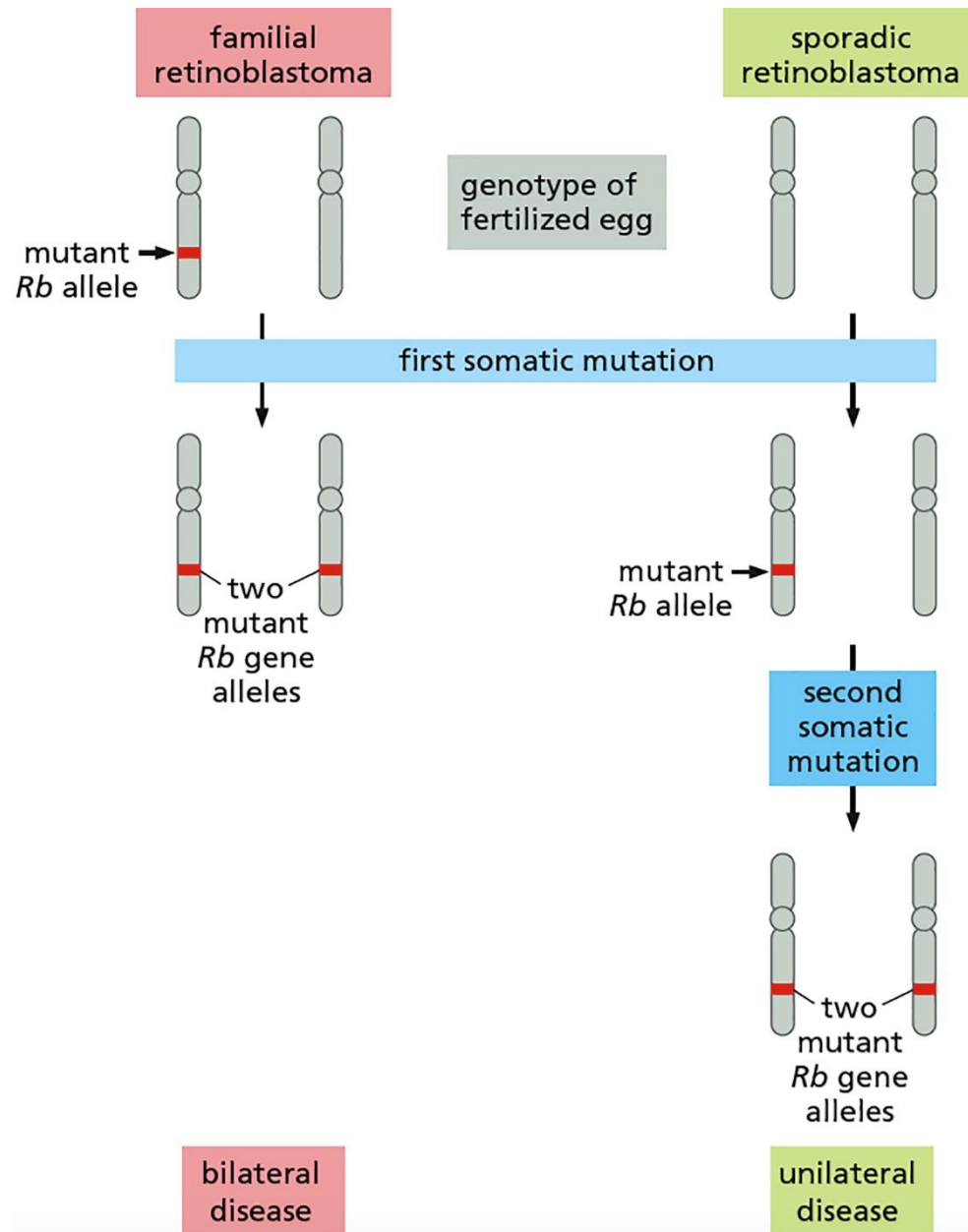
Retinoblastoma cancer

- Retinoblastoma is a rare type of eye cancer that primarily affects young children, usually before the age of 5.
- It originates in the retina, the light-sensitive lining on the inside of the eye.

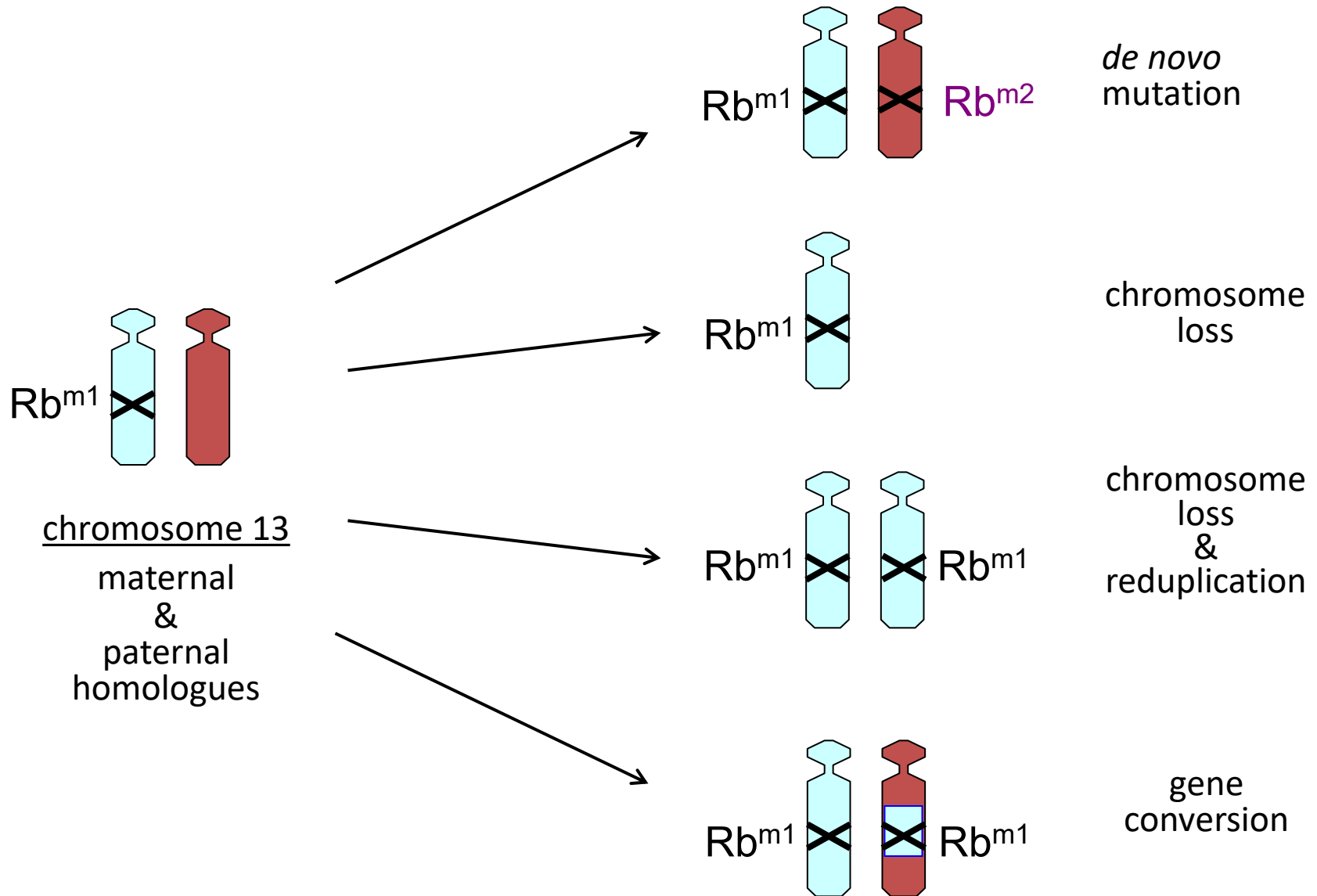


The Retinoblastoma (Rb) gene

- Cytogenetic abnormalities of chromosome 13:
 - interstitial deletions of variable length
 - always involve material from chromosome band 13q14
 - Sporadic patients: deletions in tumor cells only
 - Familial patients: deletions in both tumor & normal cells
- Is Rb susceptibility due to genetic loss at 13q14?
- If so, then the two mutations required for retinoblastoma might represent inactivation of both alleles of a single gene at 13q14

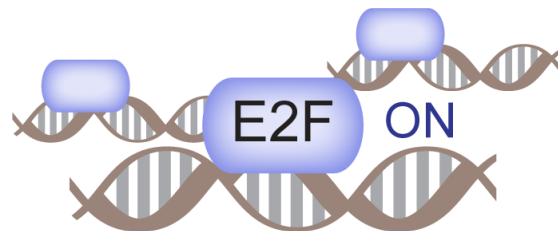


Mechanisms for inactivation of the second Rb allele in familial patients



E2F transcription factors

- The E2F family is a group of transcription factors that play an important role in regulating the cell cycle.
- G1/S transition is mediated by the E2F family of transcription factors
- E2F binds the promoters of genes required for cell cycle progression (G1/S transition and S phase).



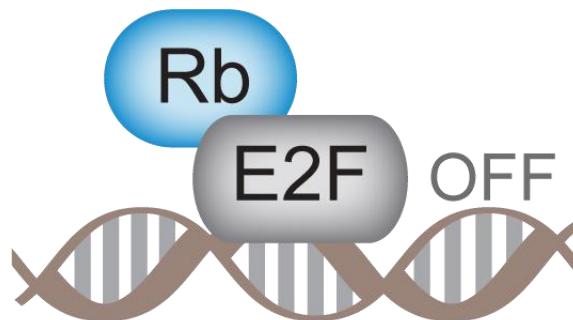
Expression of cell-cycle proteins

Some S phase genes regulated by E2F

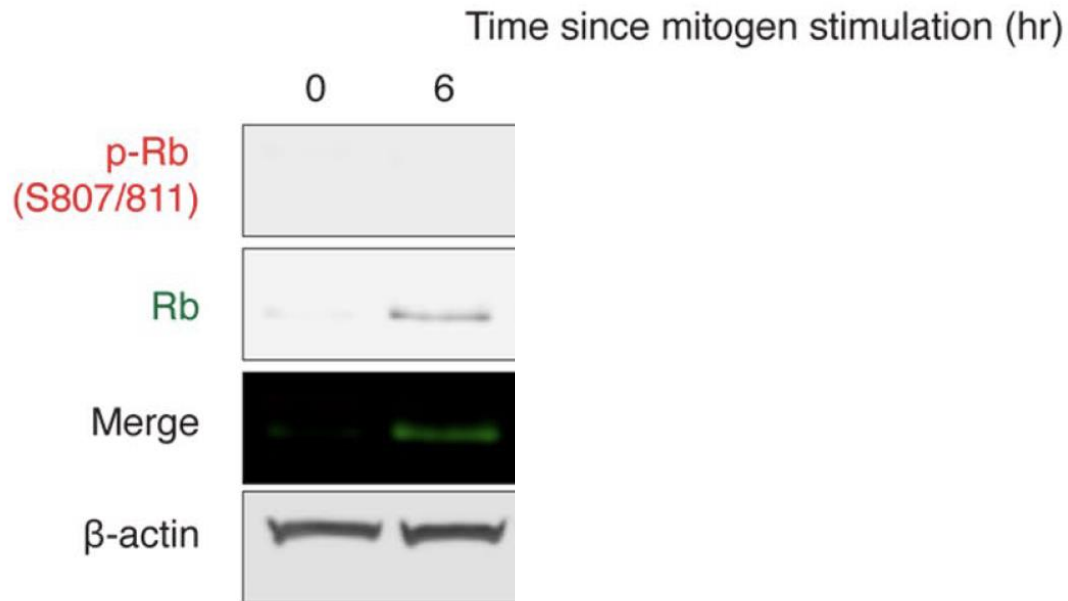
<u>S phase gene</u>	<u>Function</u>
• thymidine kinase	nucleotide synthesis
• DHFR (dihydrofolate reductase)	“ “
• DNA polymerase α	DNA replication
• ORC1	“ “
• histone H2A	chromosome assembly
• cyclin E	cell cycle progression
• cyclin A	“ “ “

Quiescent cells

- Hypophosphorylated Rb binds promoter-bound E2F
- Rb inactivates transcription by E2F
- S phase genes are repressed
- G1/S transition is blocked

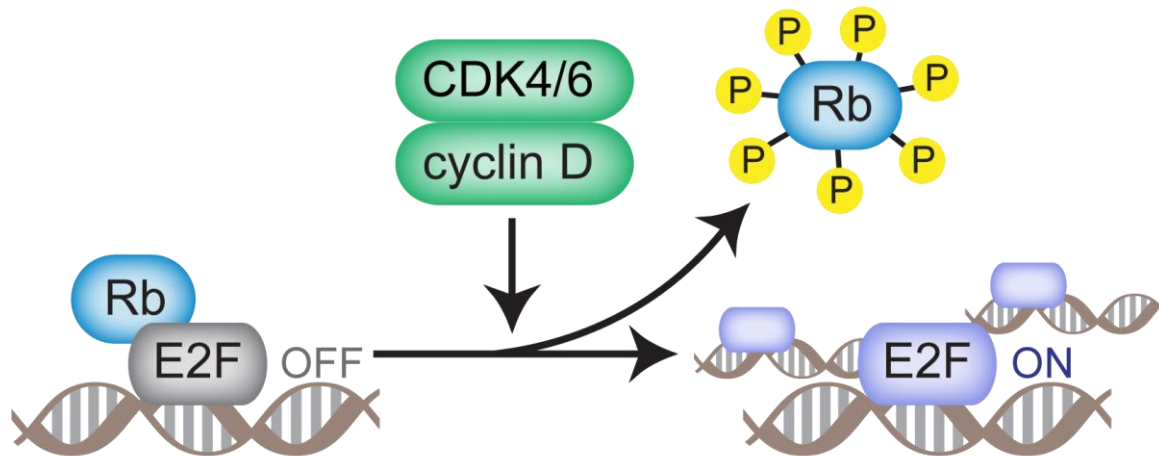


Rb is regulated by phosphorylation



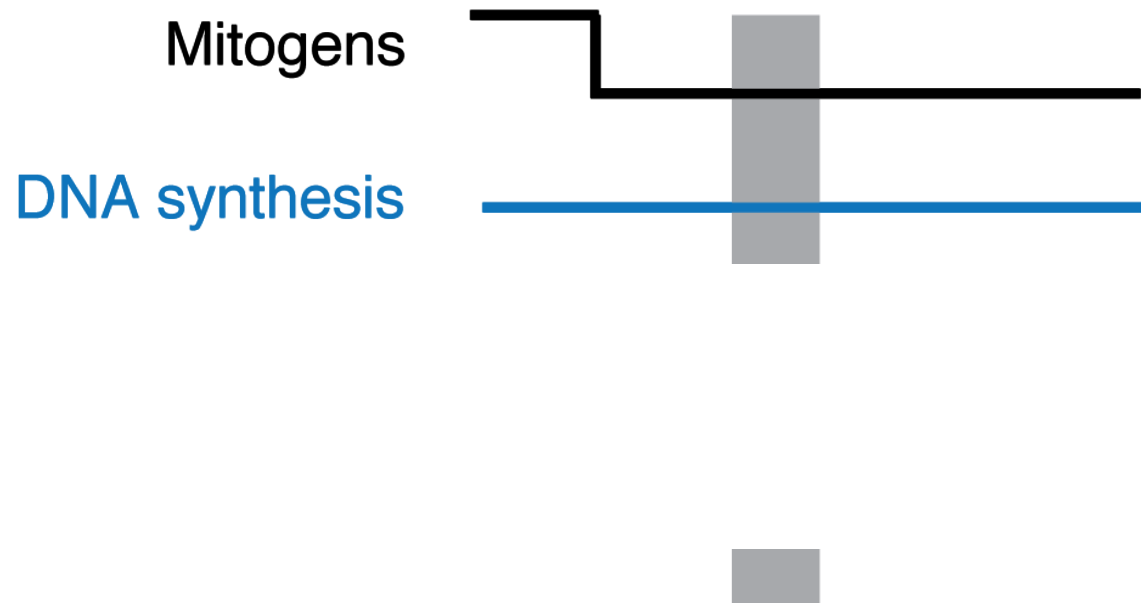
CDK4/6 phosphorylates Rb, liberating E2F

- CDK4/6-cyclin D phosphorylates Rb
- Hyperphosphorylated Rb dissociates from E2F
- E2F activates transcription of S phase genes
- Cells enter S phase

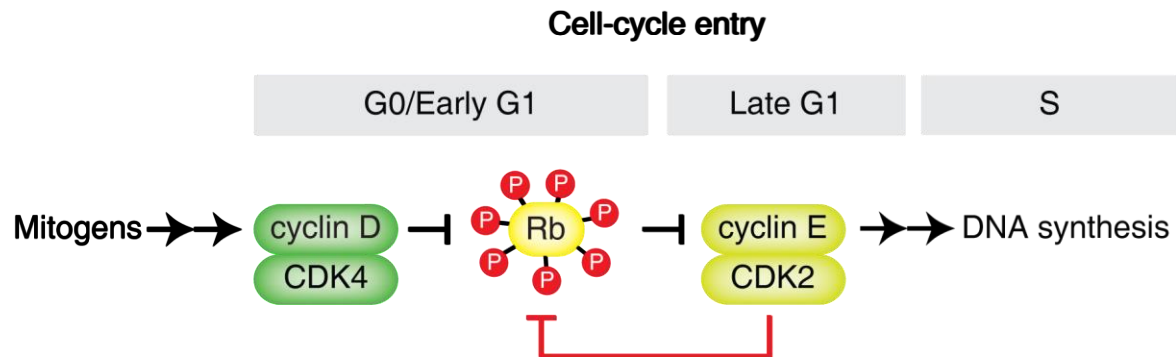
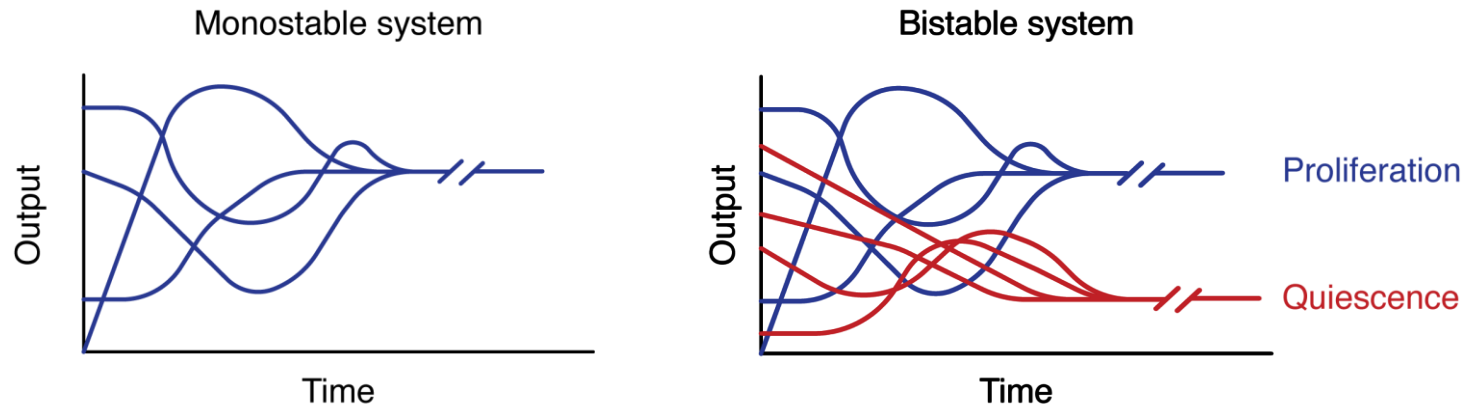


The G1/S checkpoint in cell-cycle entry

A window of mitogen requirement in G1
(The restriction point)

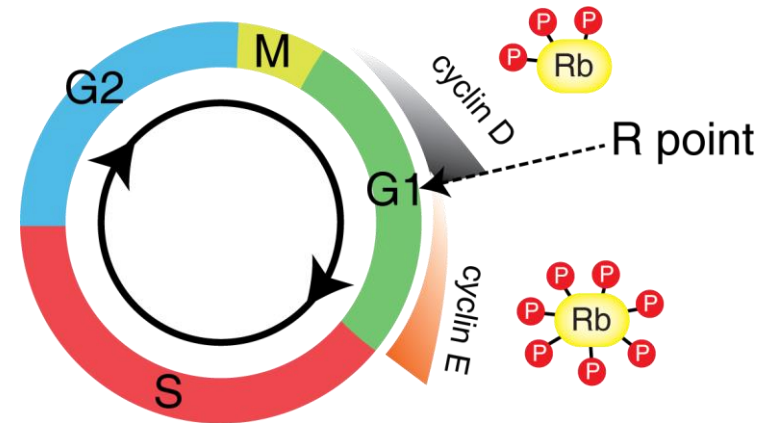
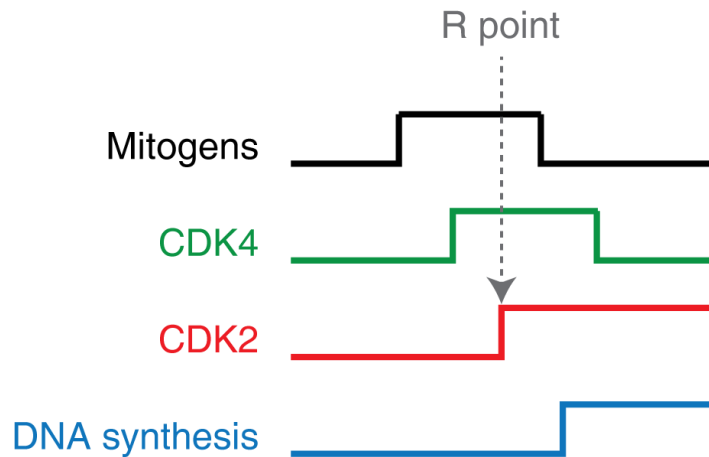


Bistability in cell-cycle entry?

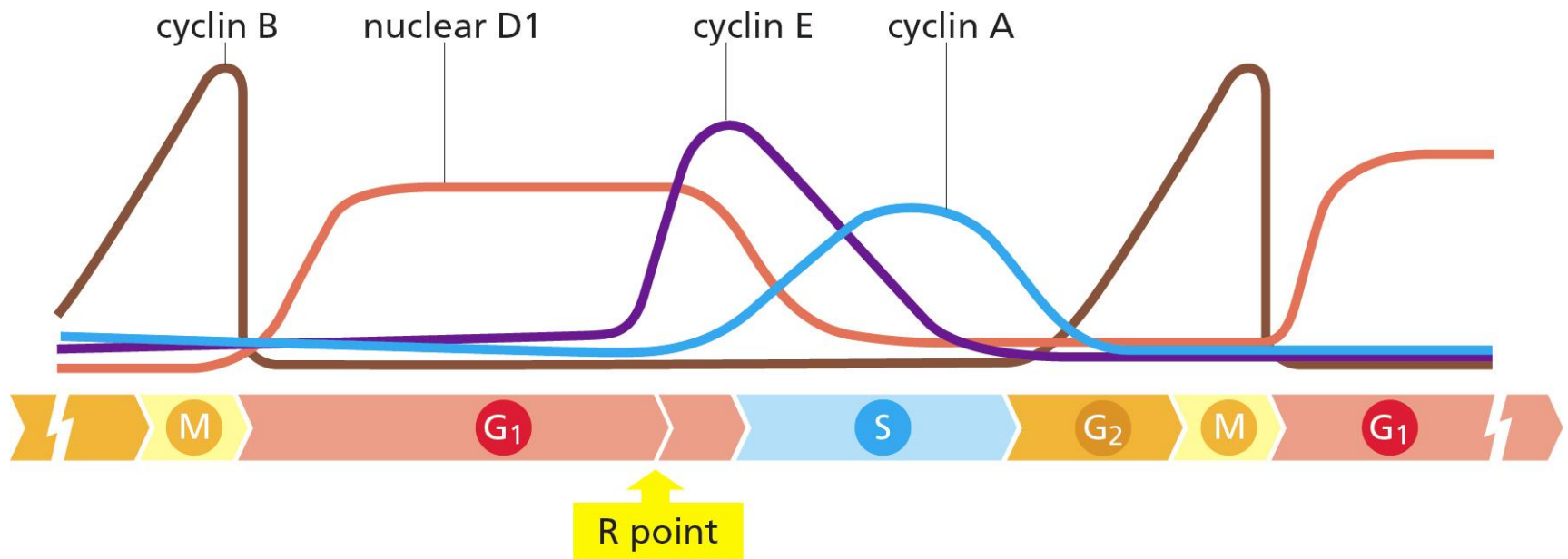


Hinds et al., 1992; Lundberg et al., 1998; Harbour et al., 1999; Yao et al., 2011;

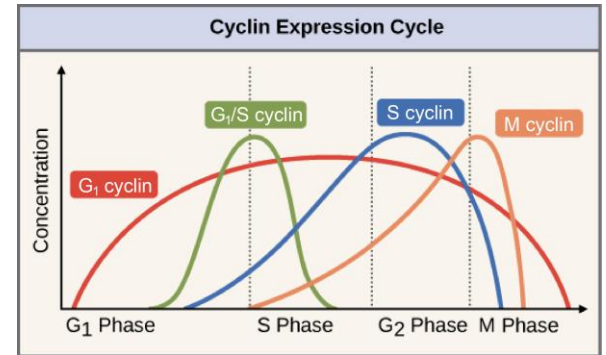
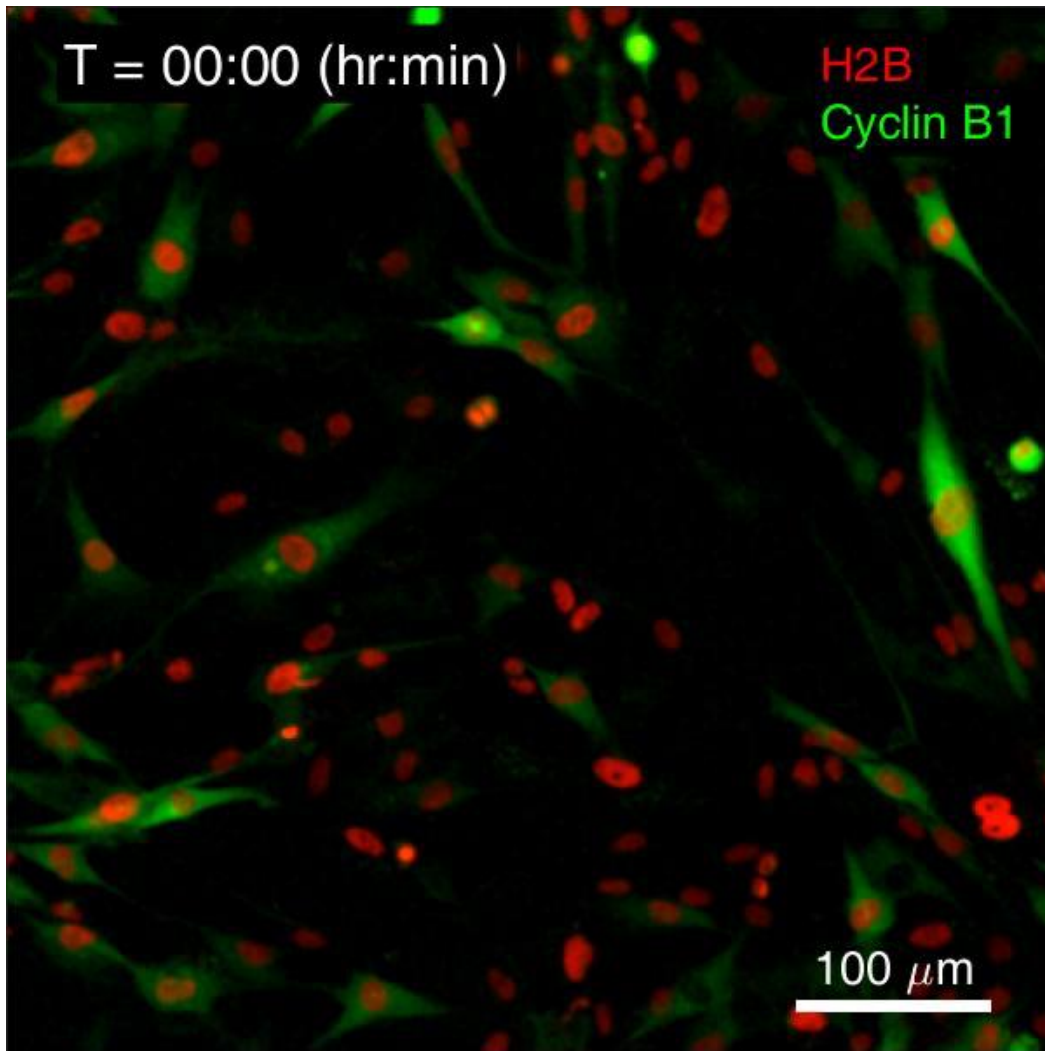
Bistability in cell-cycle entry?



Fluctuation of cyclins

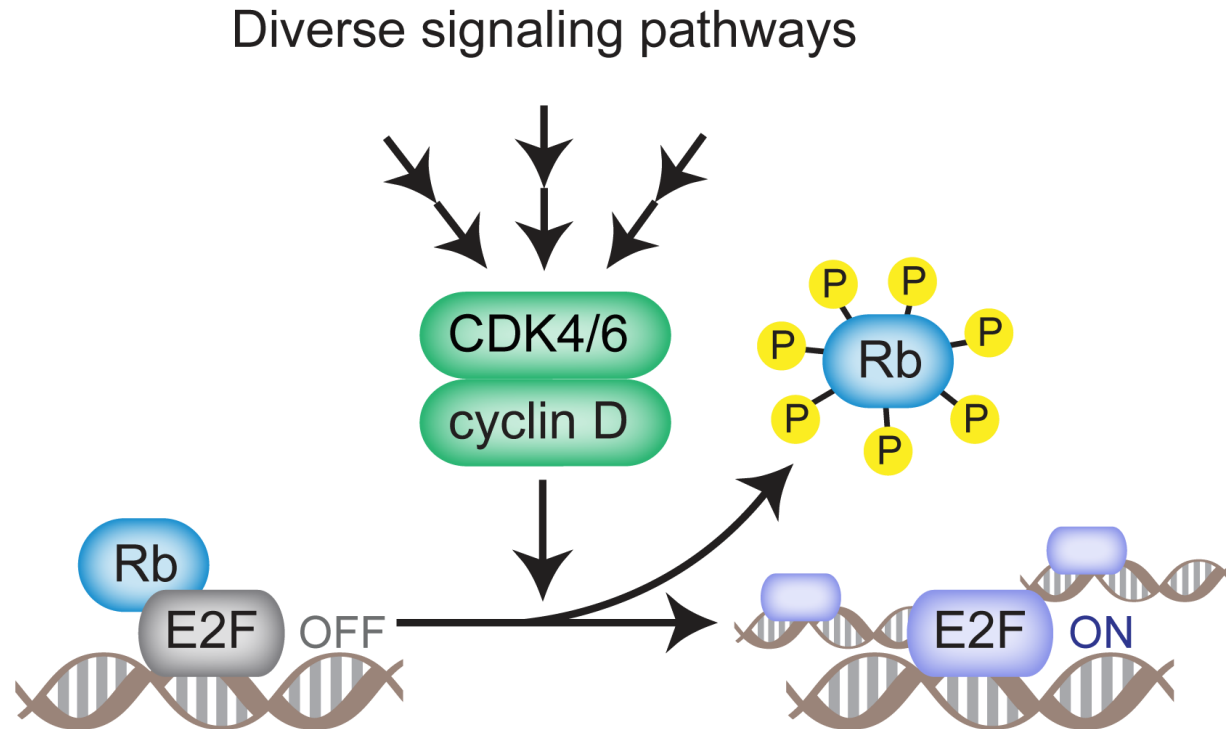


Cyclin B protein expression

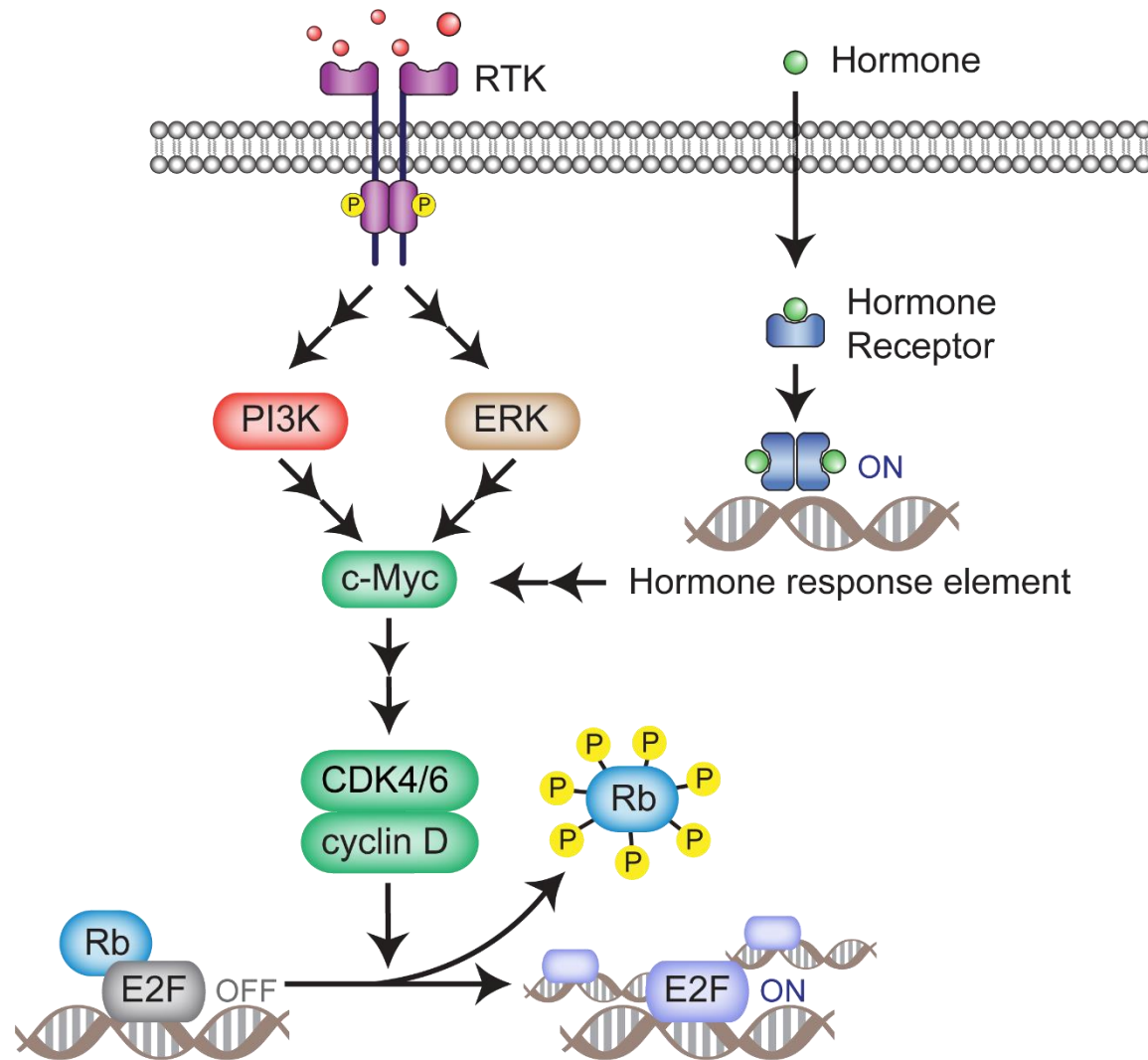


From "Control of the cell cycle: Figure 2"

Diverse signaling pathways regulate CDK4/6 activity

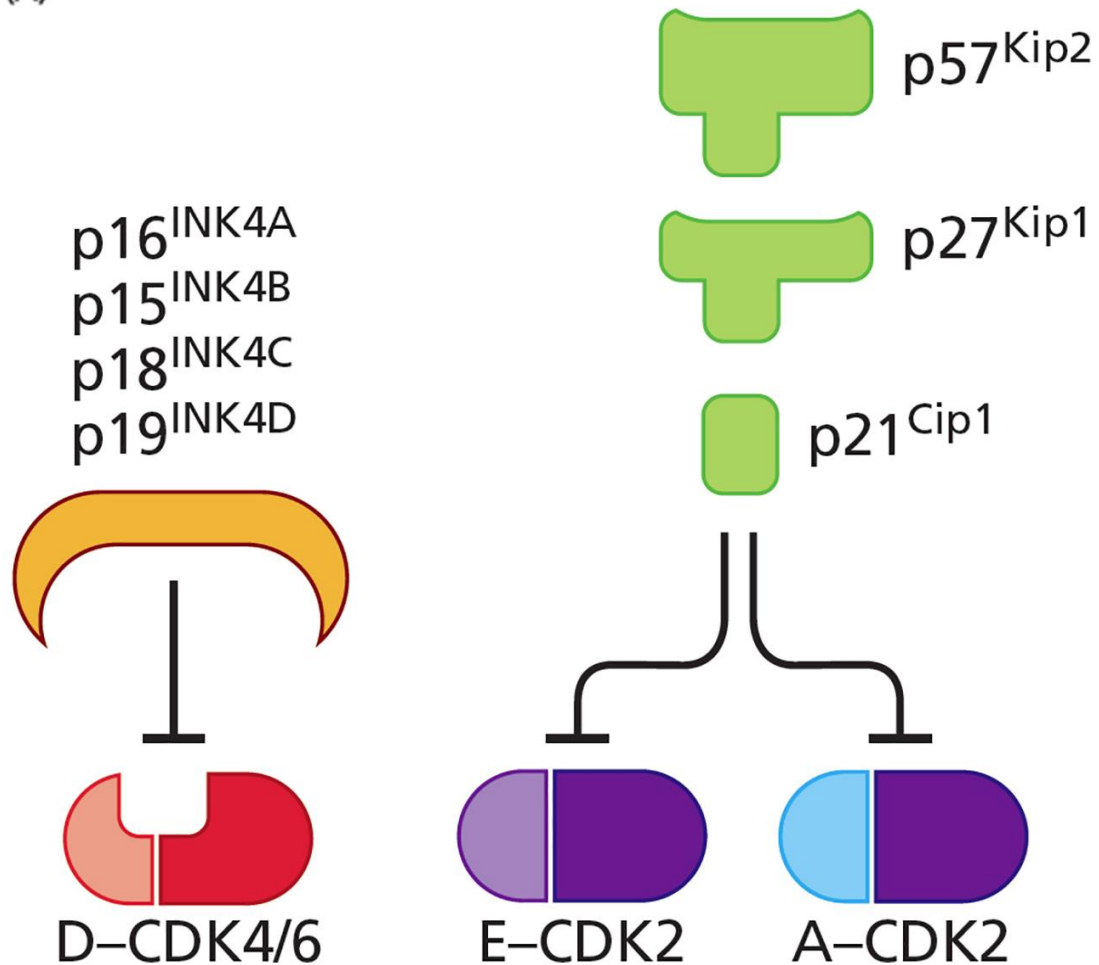


Diverse signaling pathways regulate CDK4/6 activity



CDK inhibitor proteins

(A)



CDK inhibitor proteins

INK4 (Inhibitors of CDK4) Family:

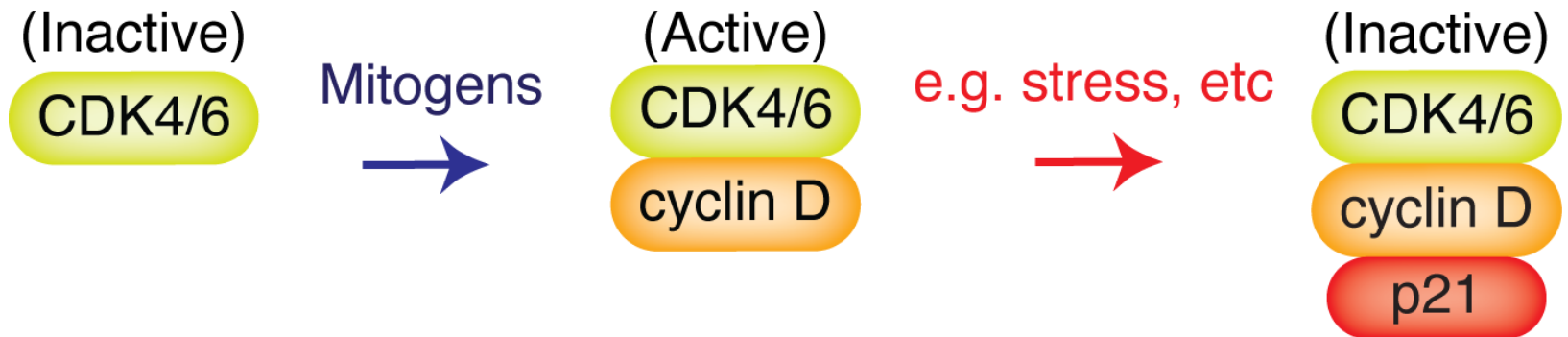
- p15, p16, p18, p19
- These proteins are composed of multiple ankyrin repeats and bind only to CDK4/6



CDK inhibitor proteins

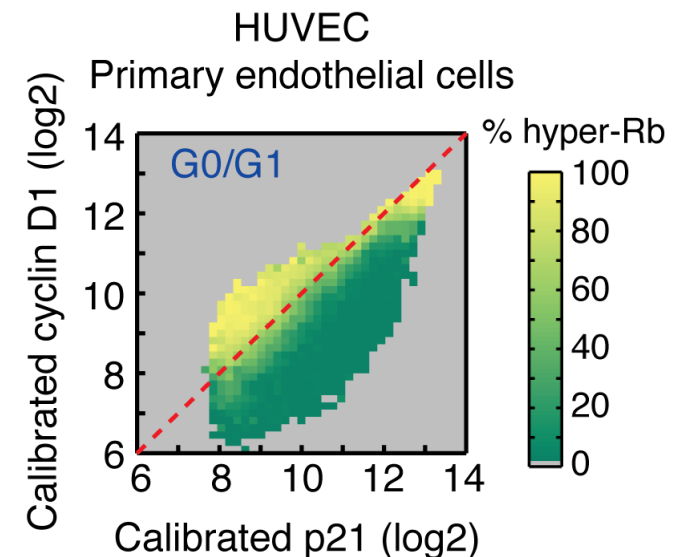
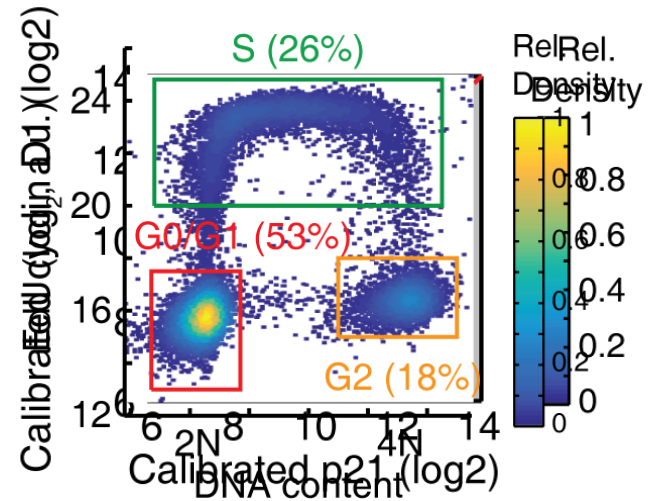
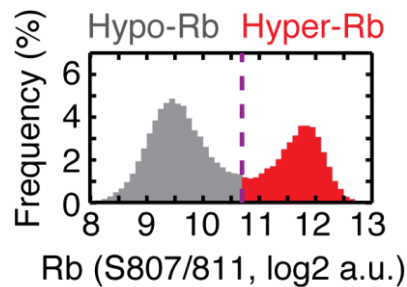
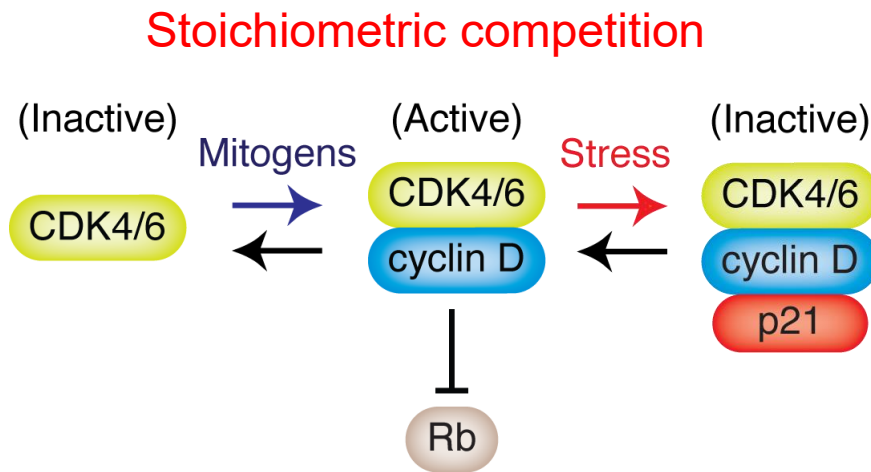
CIP/KIP (CDK interacting protein/Kinase inhibitory protein) Family:

- p21, p27, p57
- Members of the CIP/KIP family bind to and inhibit the active cyclin/CDK complex

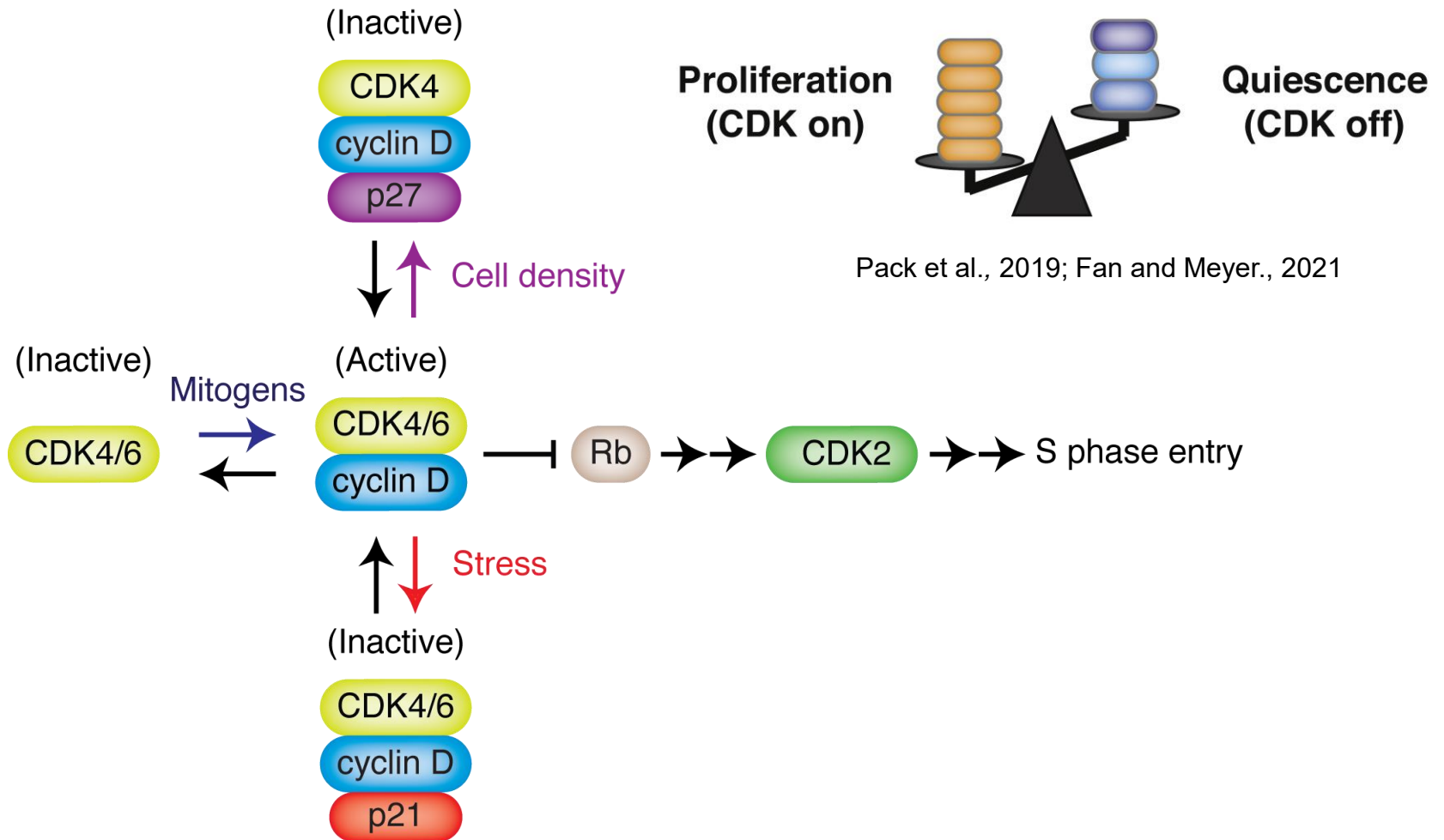


Regulation of CDK4/6 by cyclin D and p21

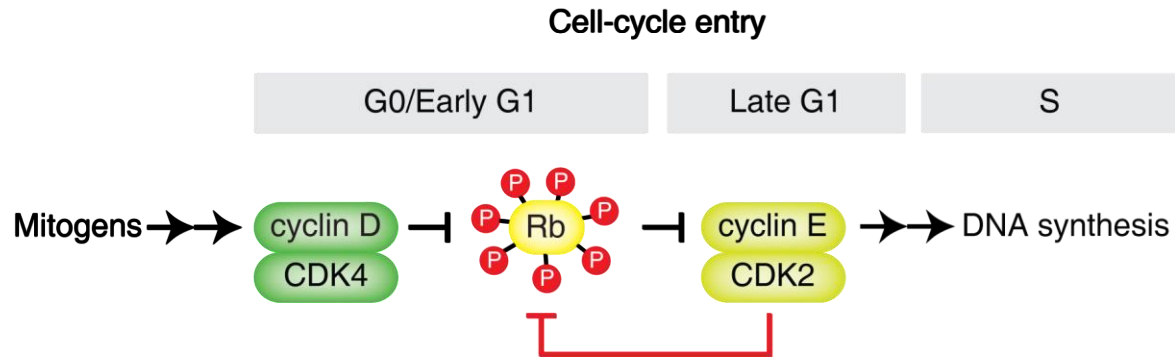
Yang et al. 2017



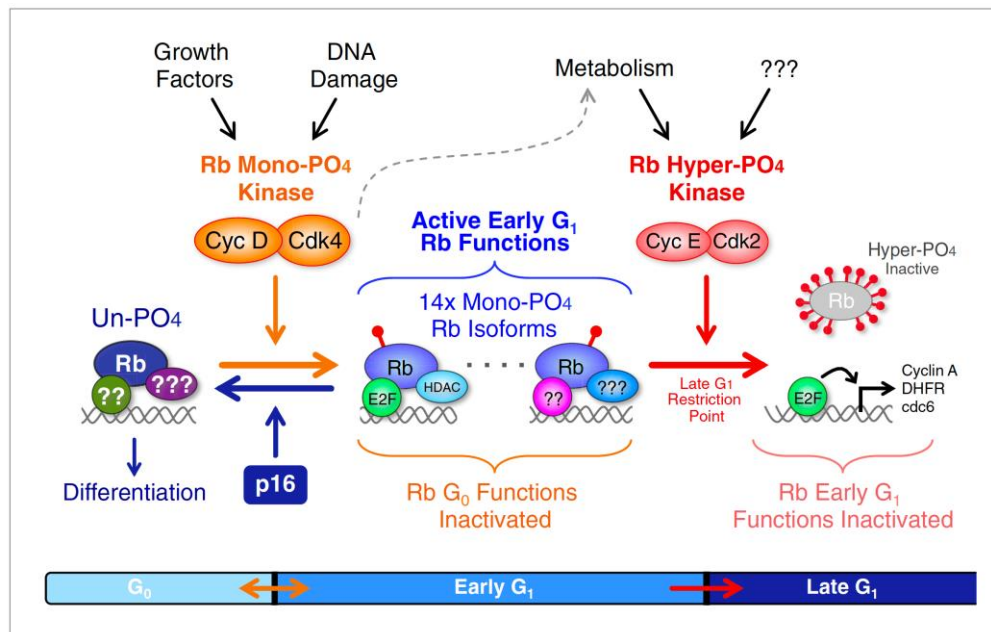
Competing mitogen and stress signaling control cell-cycle entry



The role of CDK4/6 in cell-cycle entry



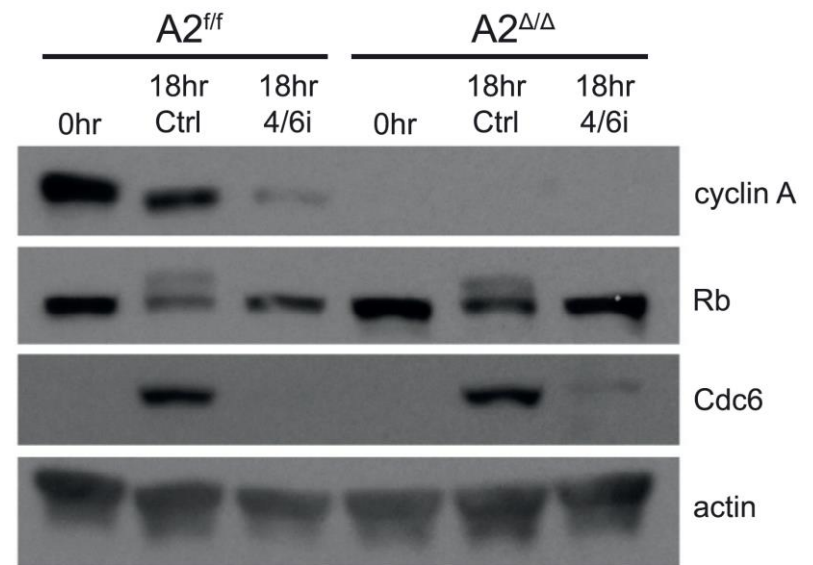
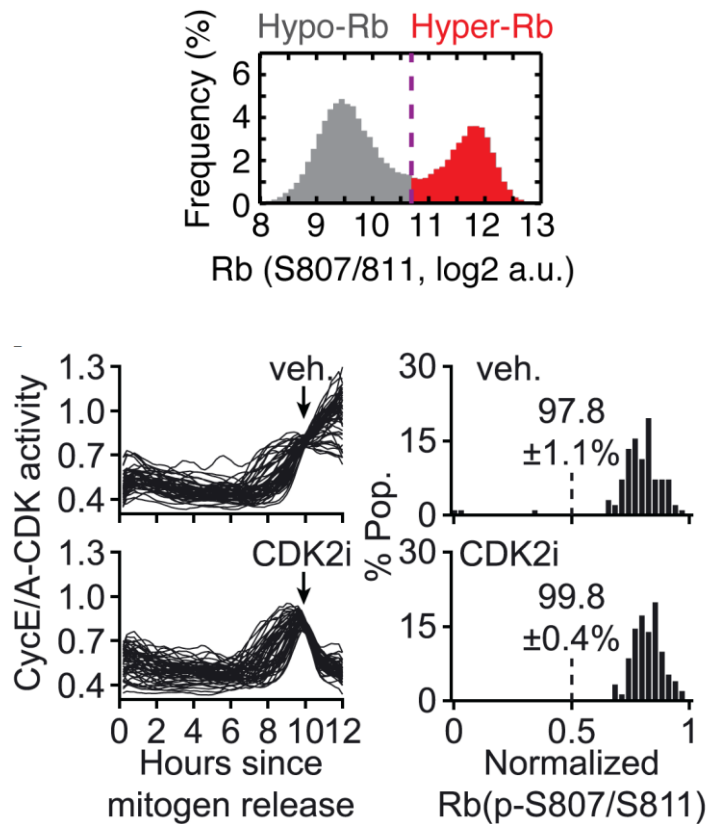
Hinds et al., 1992; Lundberg et al., 1998; Harbour et al., 1999; Yao et al., 2011;



Narasimha et al., 2014

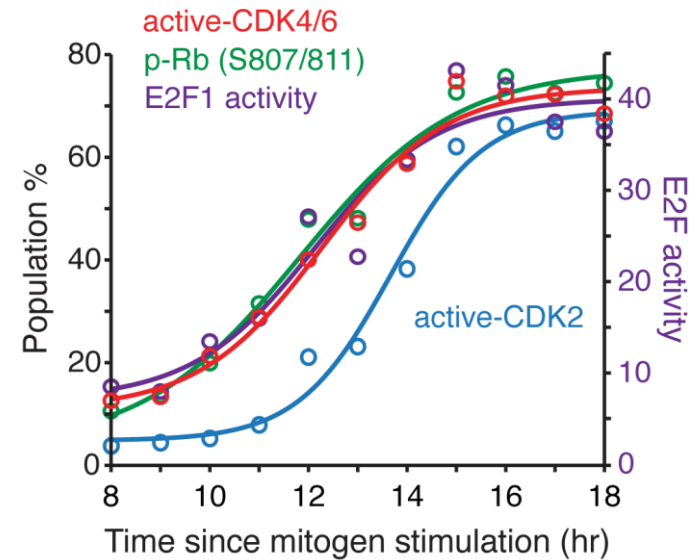
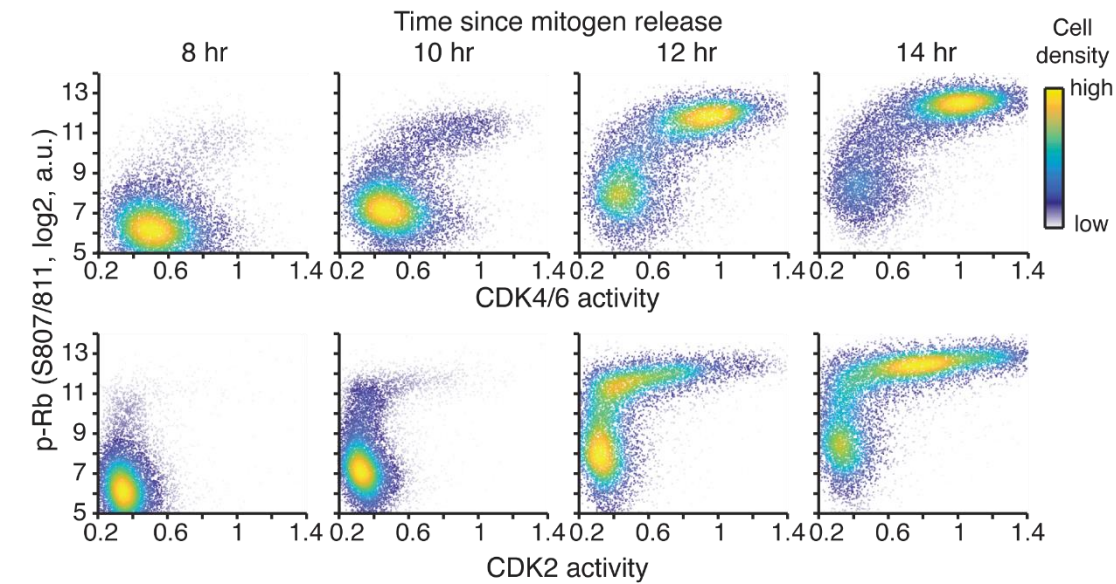
The role of CDK4/6 in cell-cycle entry

Chung et al., 2019



The role of CDK4/6 in cell-cycle entry

Kim et al., 2022



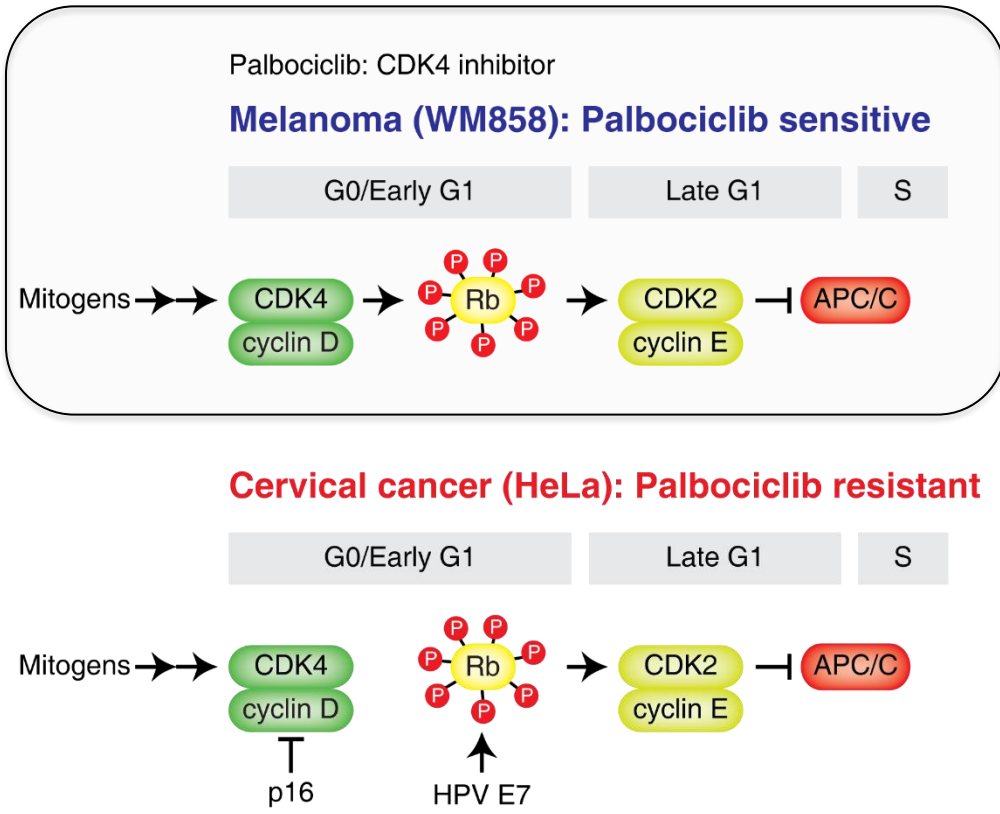
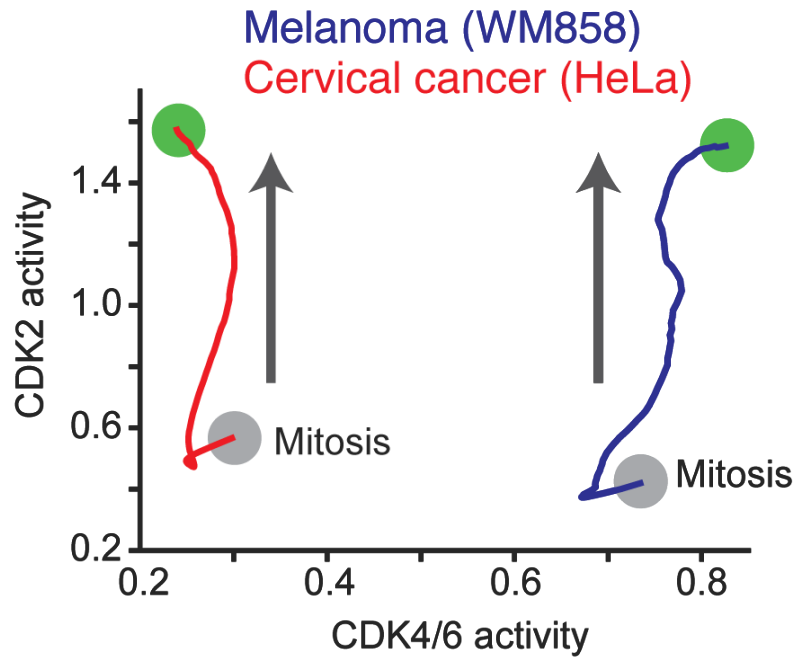
Direct inactivation of Rb in tumors

- Rb gene deletion (occurs in retinoblastoma)
- point mutations in the Rb pocket (in retinoblastoma)
- occupancy of the Rb pocket by early proteins of DNA tumor viruses
 - human papilloma virus (HPV), the main etiological agent of human cervical carcinomas
 - HPV encodes two proteins required for tumorigenesis
 - E7 binds the pocket of hypophosphorylated Rb
 - Deregulation of E2F (and the G1/S transition)

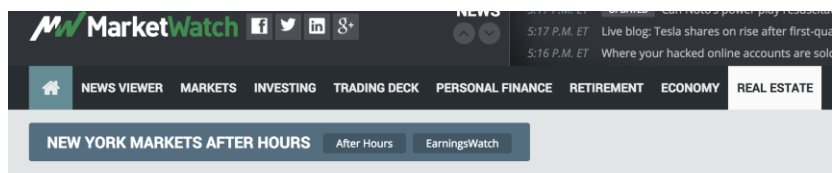
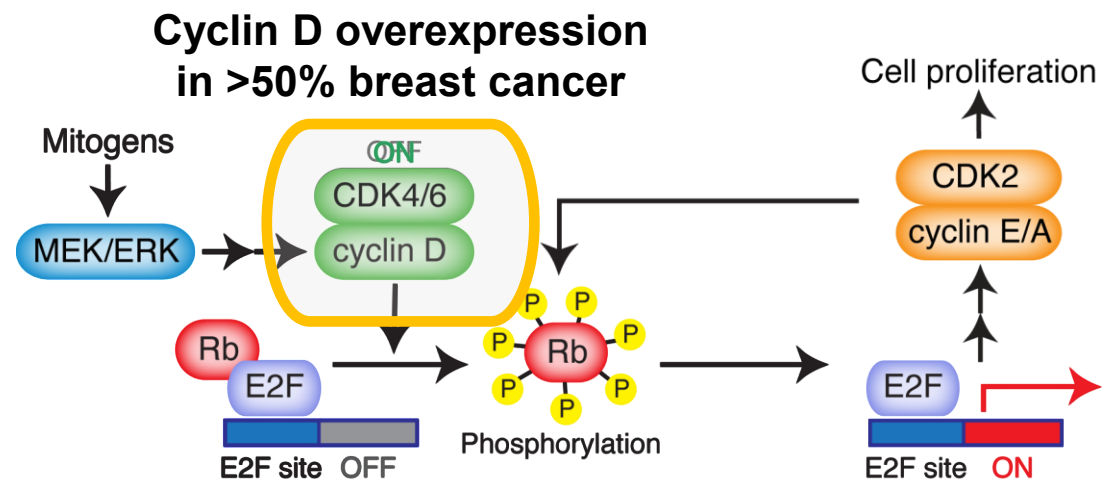
Indirect inactivation of Rb in tumors

- overexpression of cyclin D1
 - breast cancer, B cell lymphoma
 - loss of p16, an inhibitor of Cdk4
 - many human cancers
 - inherited point mutation in Cdk4 that renders it insensitive to inhibition by p16
 - familial melanoma
- » Inactivation of the Rb pathway occurs in many human tumors!

Rb-dependent and -independent cell-cycle entry



CDK4/6 is a promising target in breast cancer

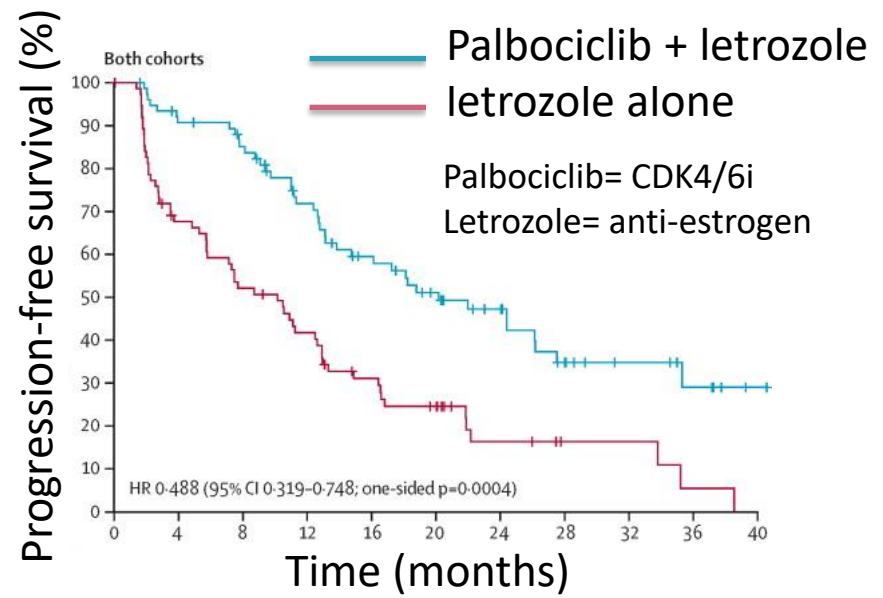


PRESS RELEASE

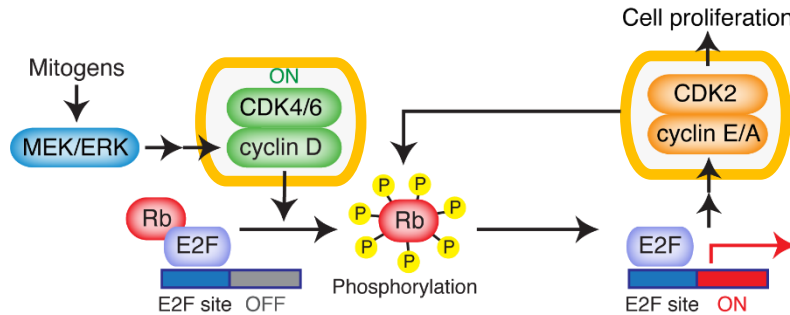
Pfizer Announces PALOMA-3 Trial For IBRANCE® (Palbociclib) Stopped Early Due To Efficacy Seen In Patients With HR+, HER2-Metastatic Breast Cancer Whose Disease Has Progressed Following Endocrine Therapy

Published: Apr 15, 2015 8:00 a.m. ET

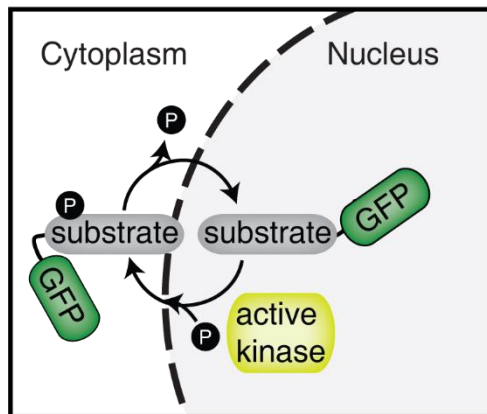
Finn et al., 2015



Live-cell sensor to monitor individual cell proliferation



Kinase Translocation Reporters (KTRs)



$$\text{Kinase activity} = \frac{\text{Cytoplasm}}{\text{Nucleus}}$$

Hahn et al., 2009 Regot et al., 2014

Spencer et al., 2013

Yang*, Cappell*, Jaimovich* et al., 2020

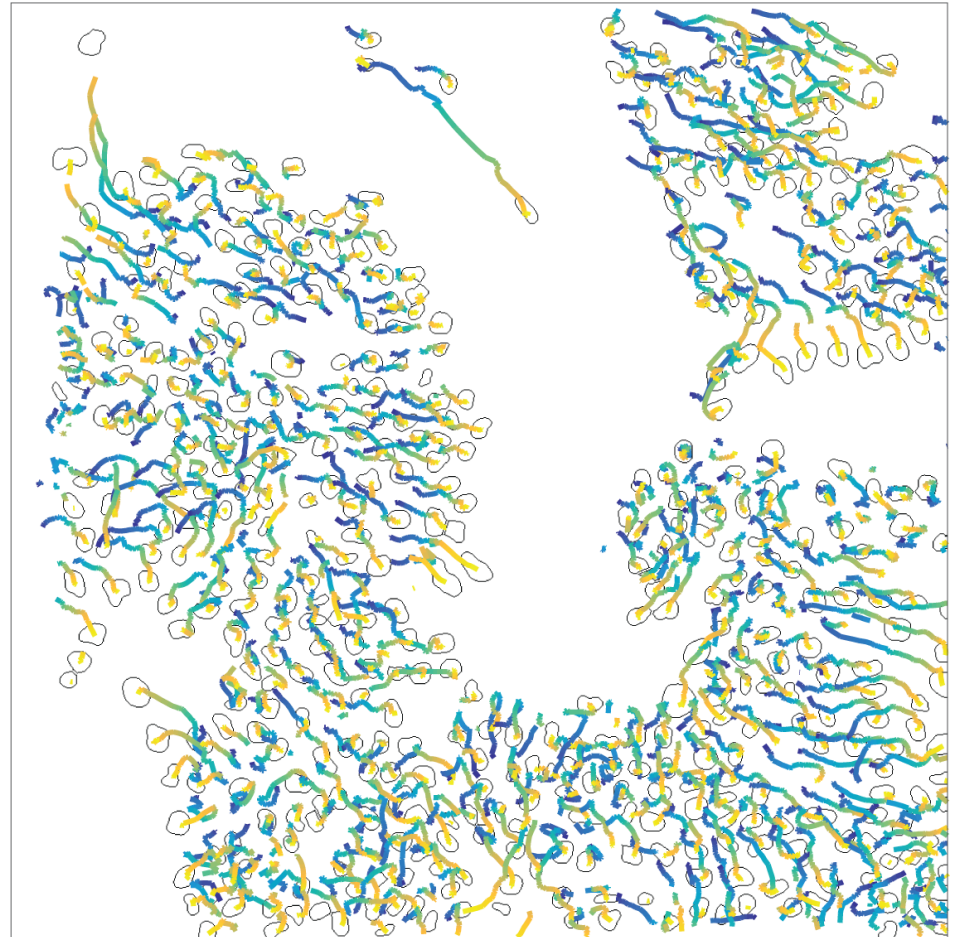
Individual cell tracking

Live cell imaging in
Highthroughput manner



Cell tracking & Classification of cell behavior

Time
(hr) 0 5 10



Drug-tolerant persister cells

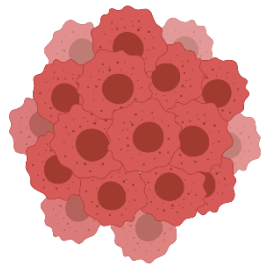


Proliferation

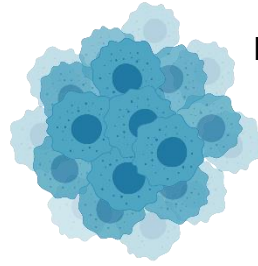


Quiescence

Drug-naïve cells



Drugs



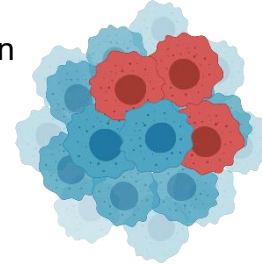
Rapid adaptation
(non-genetic)



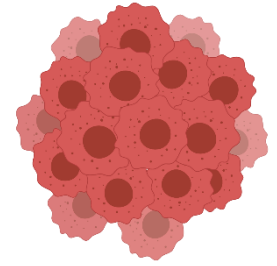
Reversible



**Drug-tolerant
persister cells**



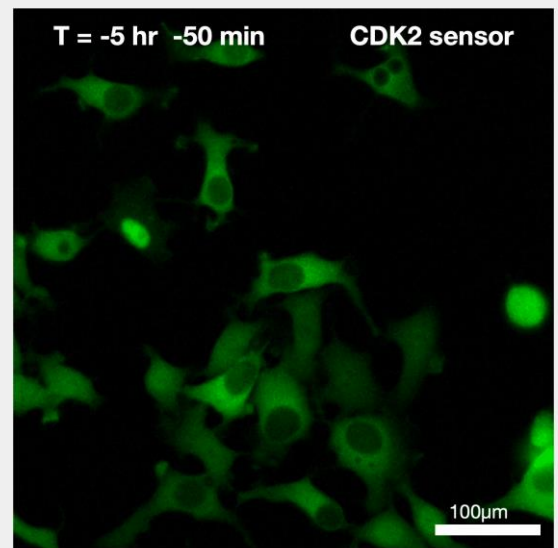
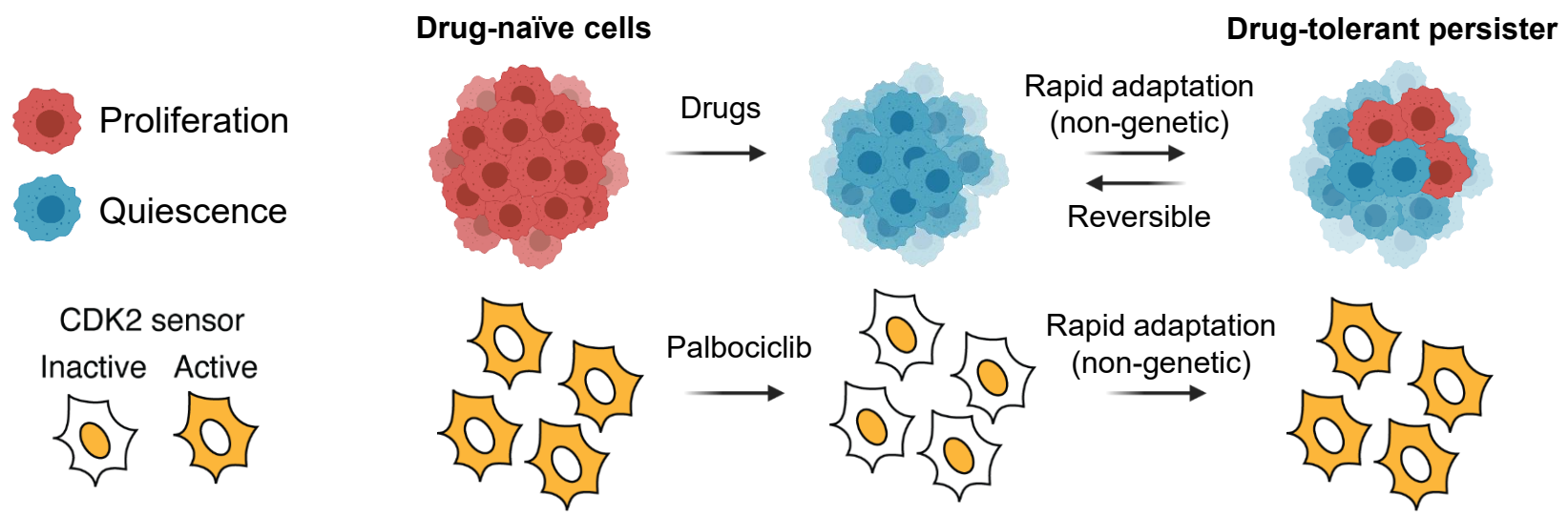
Resistant clones



Reversible

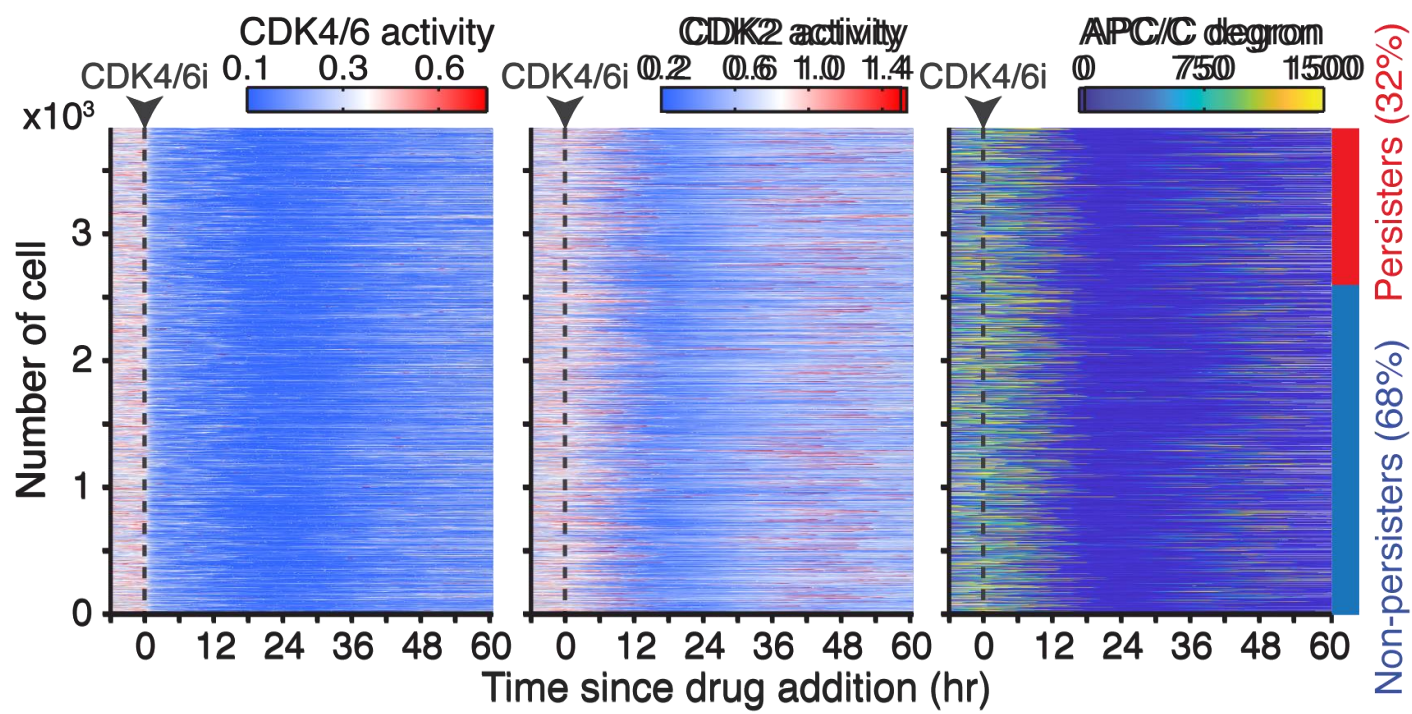
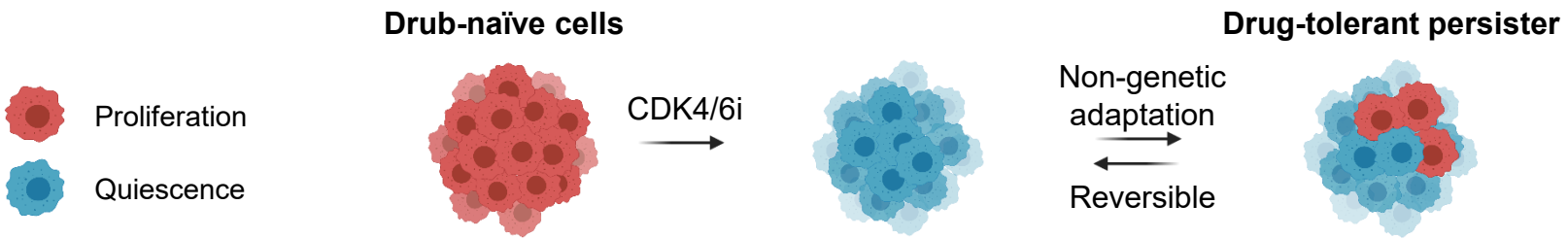


Drug-tolerant persister cells



HR⁺/HER2⁻ MCF7 cells

Drug-tolerant persister cells

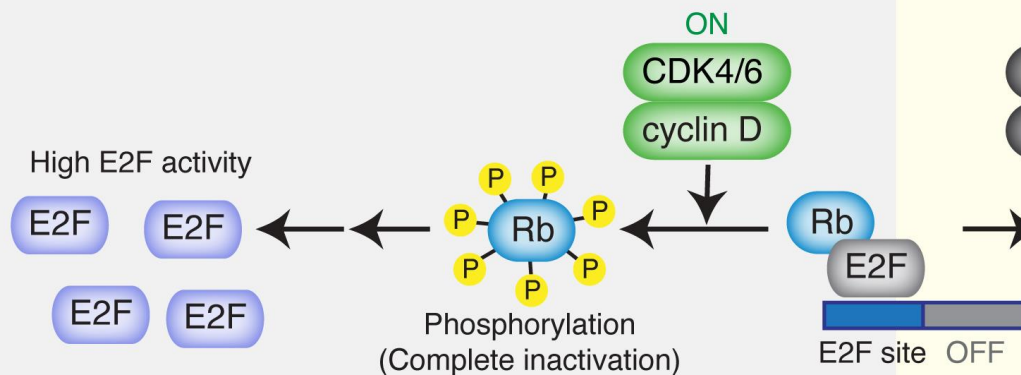


1% – 30%

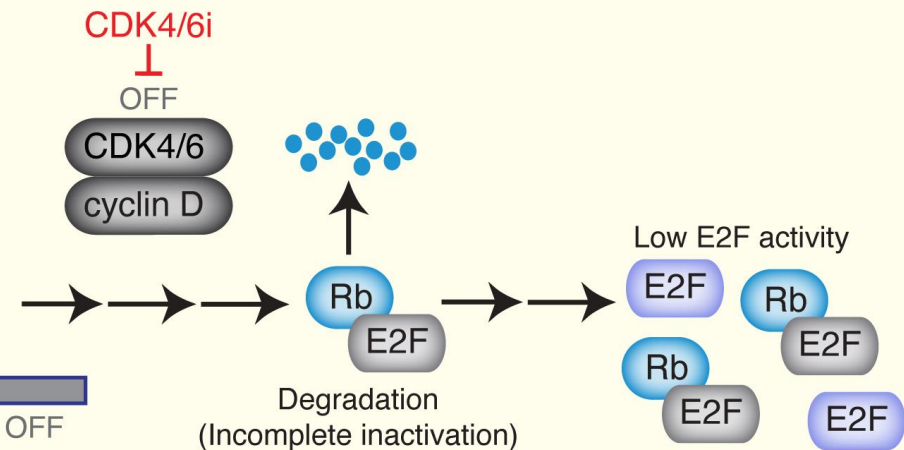
Alternative pathway for Rb inactivation

Kim et al. 2023; Zhang* and Kim* et al. 2023

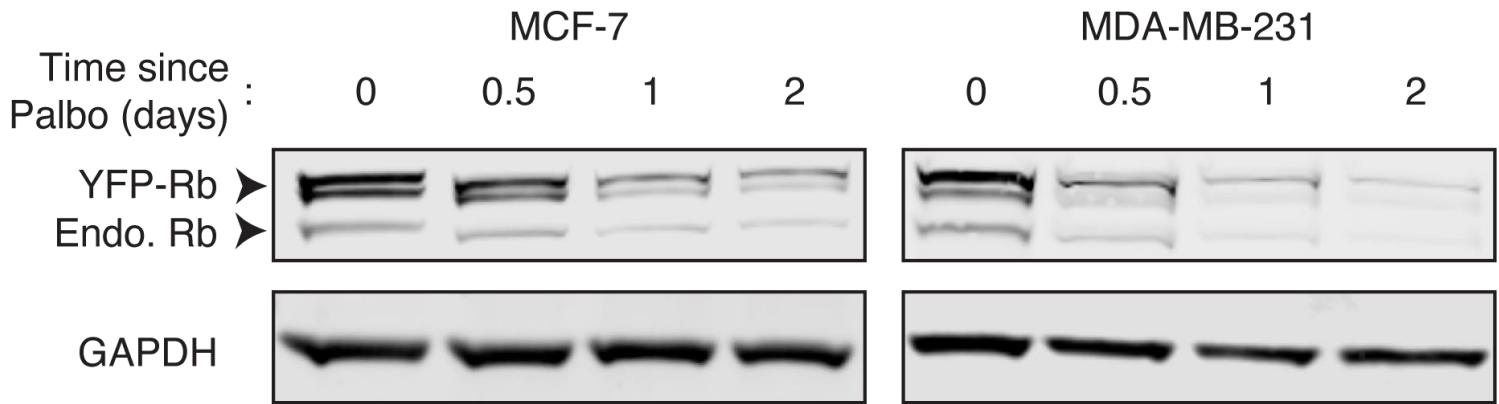
Canonical pathway



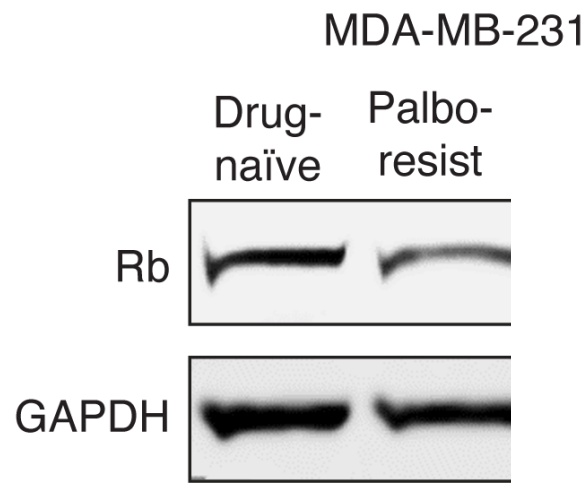
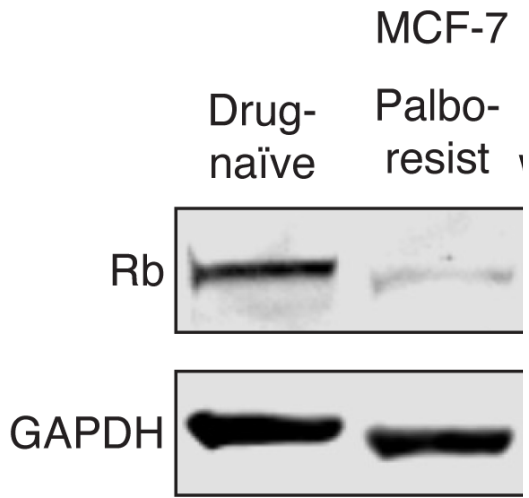
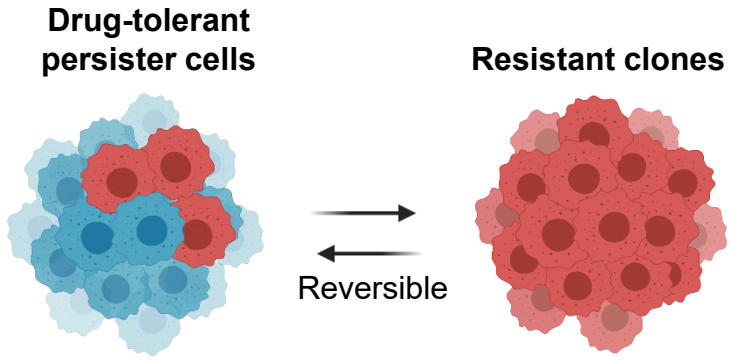
Non-canonical pathway



CDK4/6 inhibitor treatment results in Rb-protein reductions

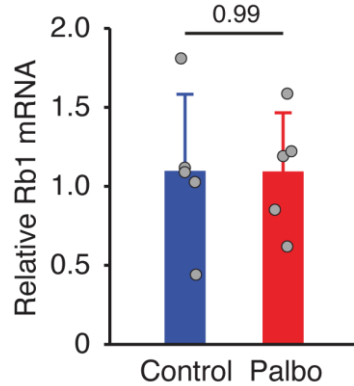
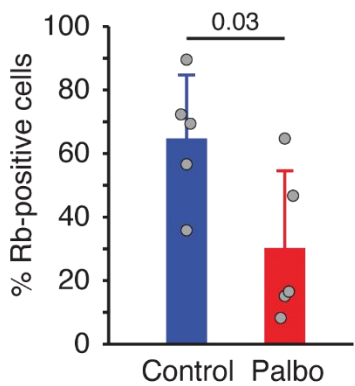
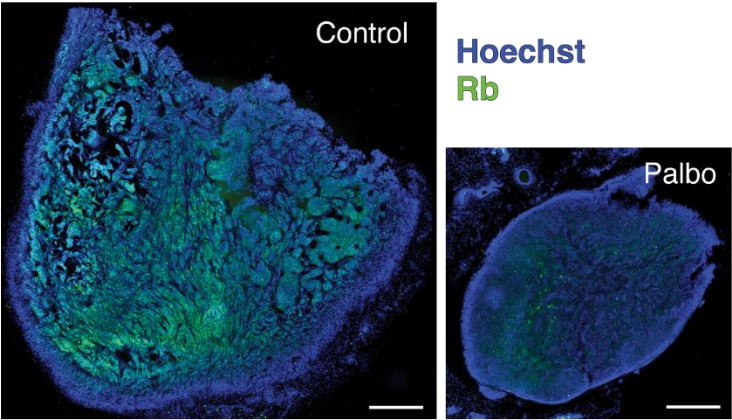
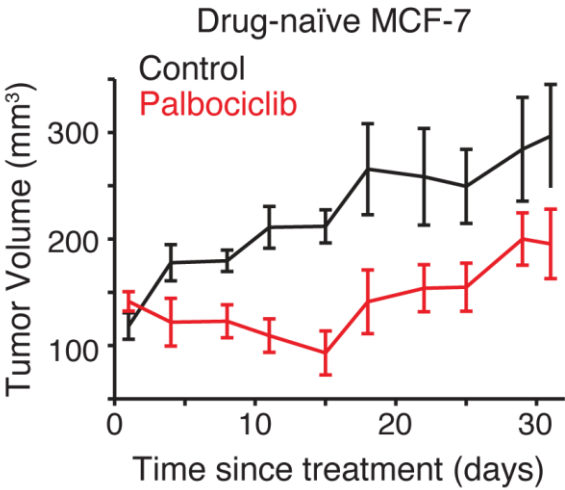
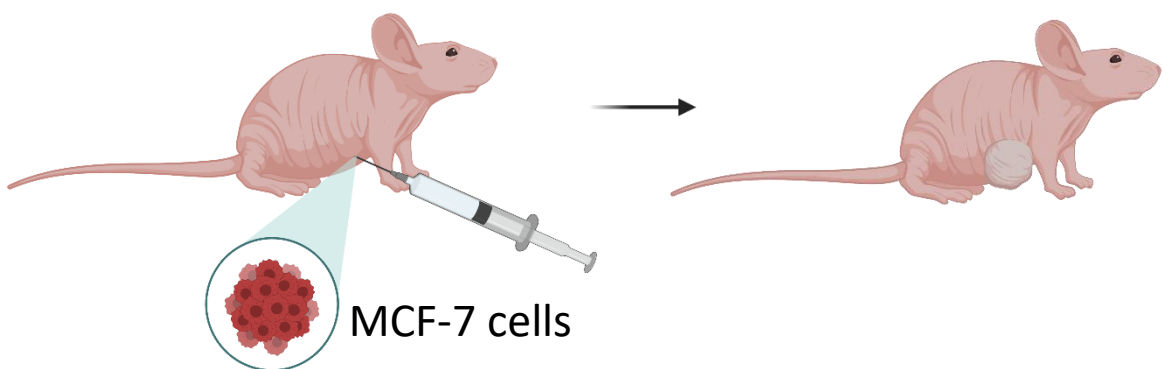


Rb-protein reductions are reversible in established CDK4/6i resistance

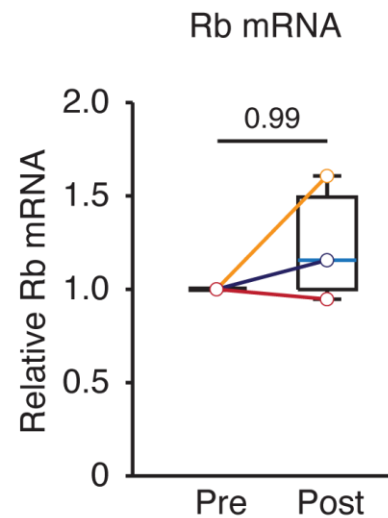
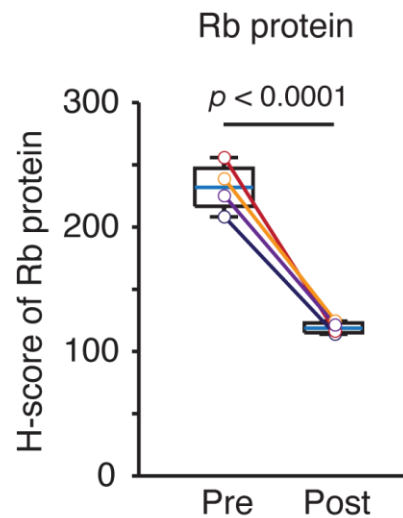
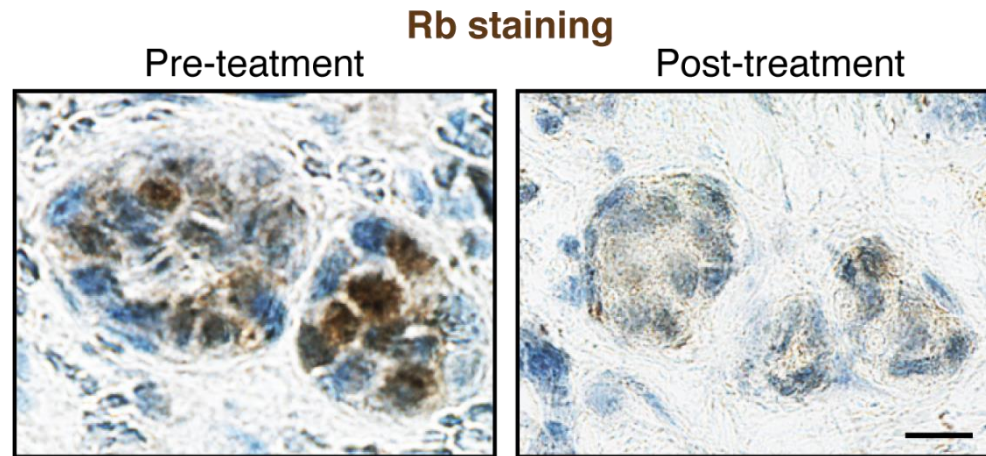


Reduced Rb protein levels in a breast cancer mouse model

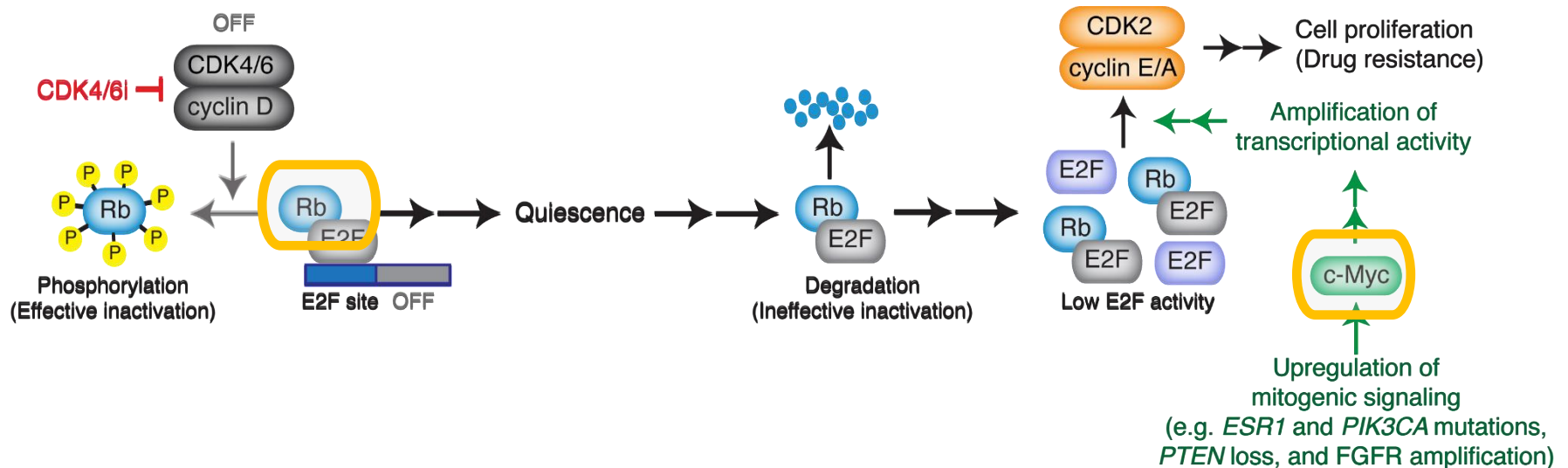
Minah Kim (CUIMC)



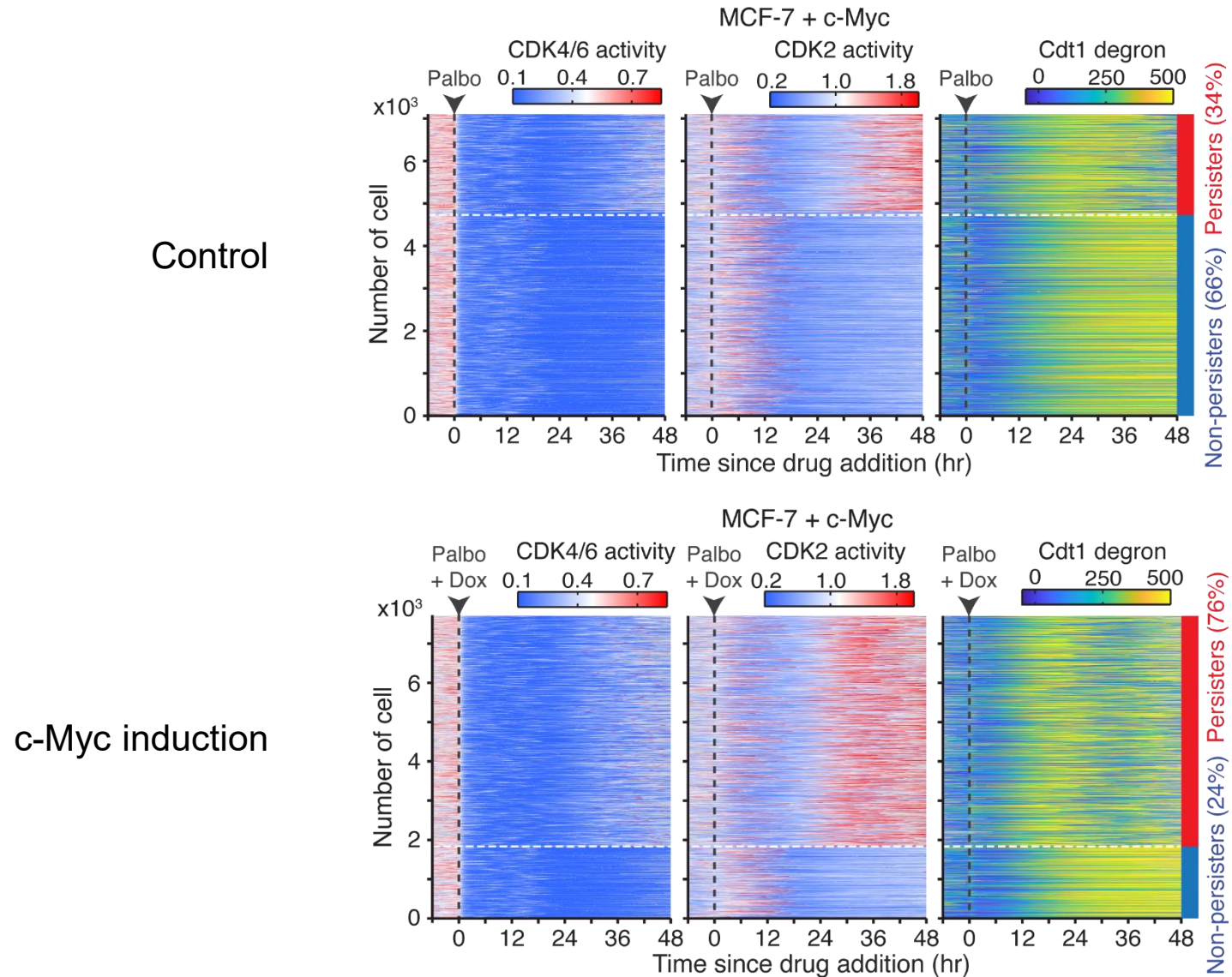
Rb-protein reduction in patient samples



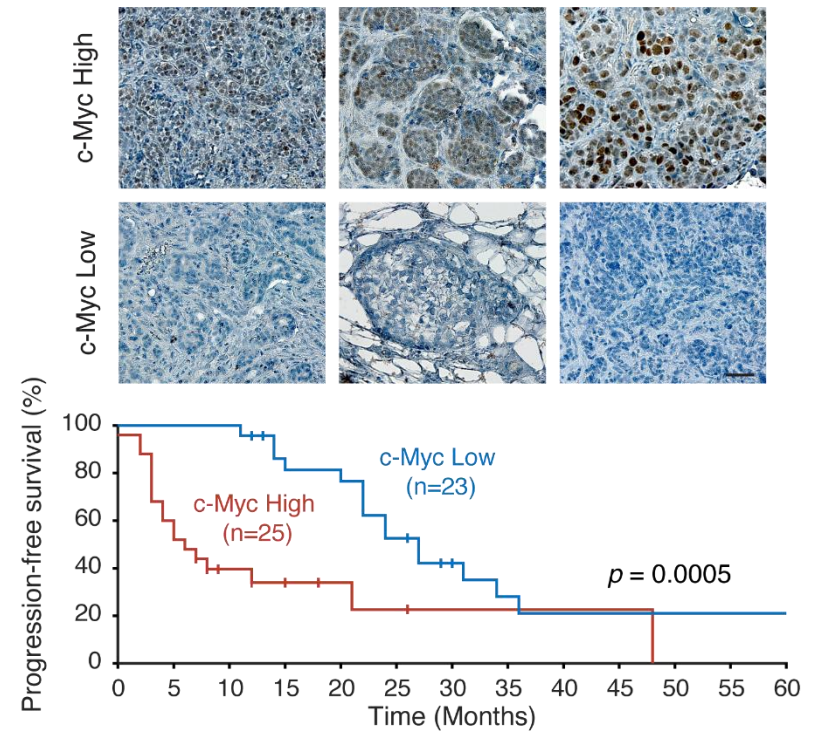
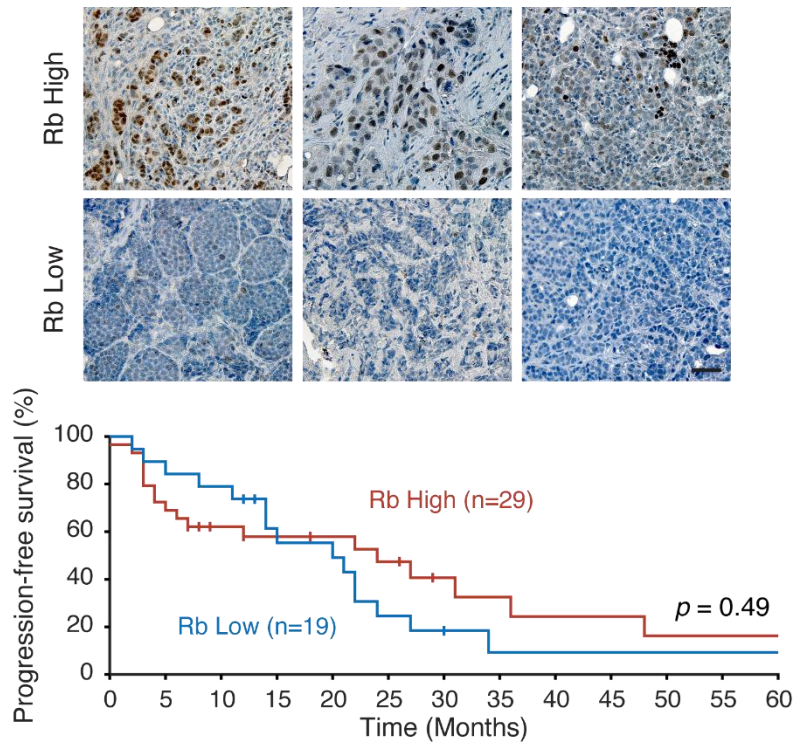
Sequential regulation of E2F activation



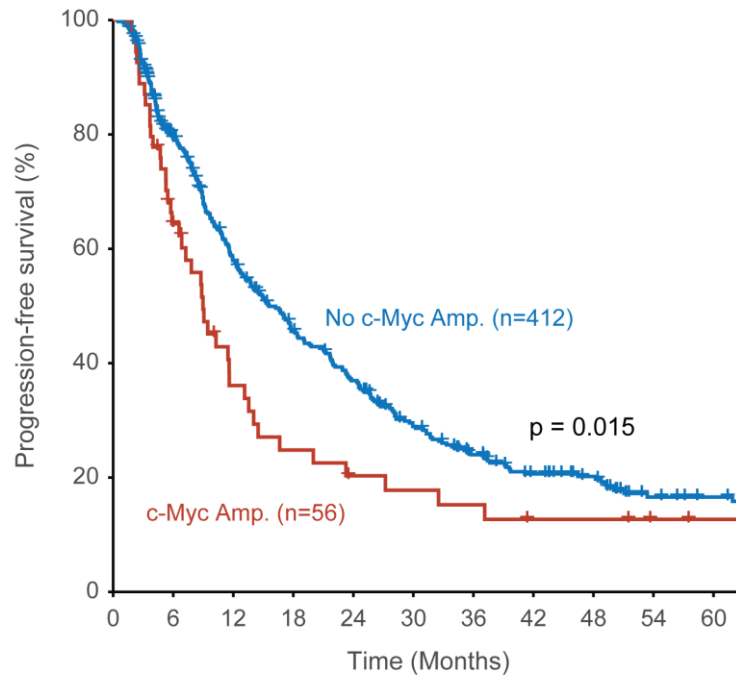
Induction of c-Myc expression facilitates CDK4/6i-tolerant persisters



Rb and c-Myc levels in pre-treatment samples and PFS



c-Myc amplification status in pre-treatment samples and PFS



Adjusted by hormone therapy alone

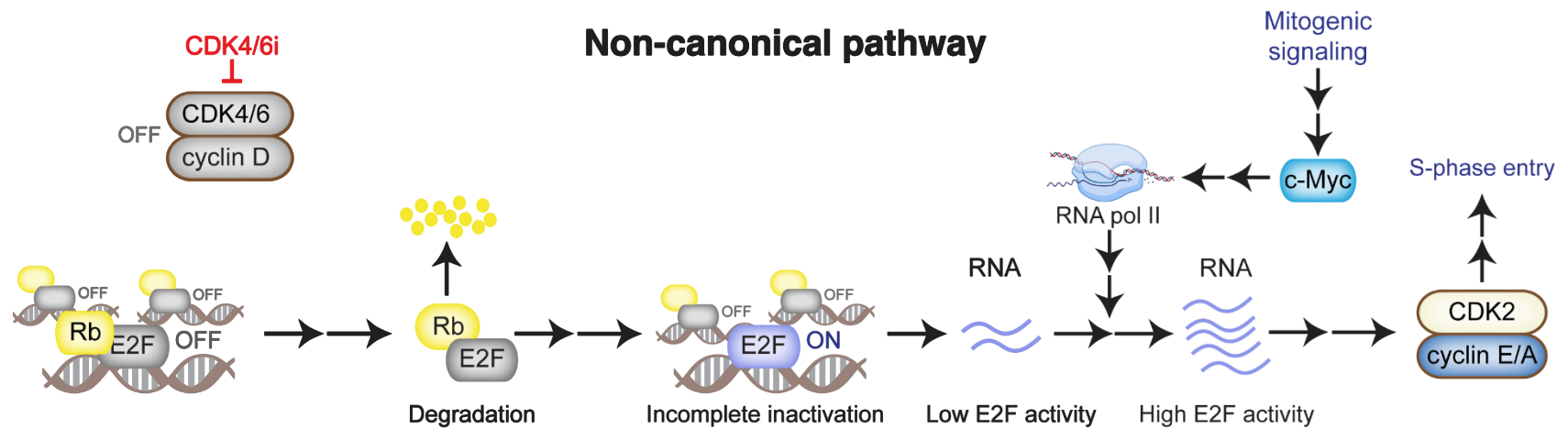
Sarat Chandarlapaty (MSKCC)

Pedram Razavi (MSKCC)

Anton Safonov (MSKCC)

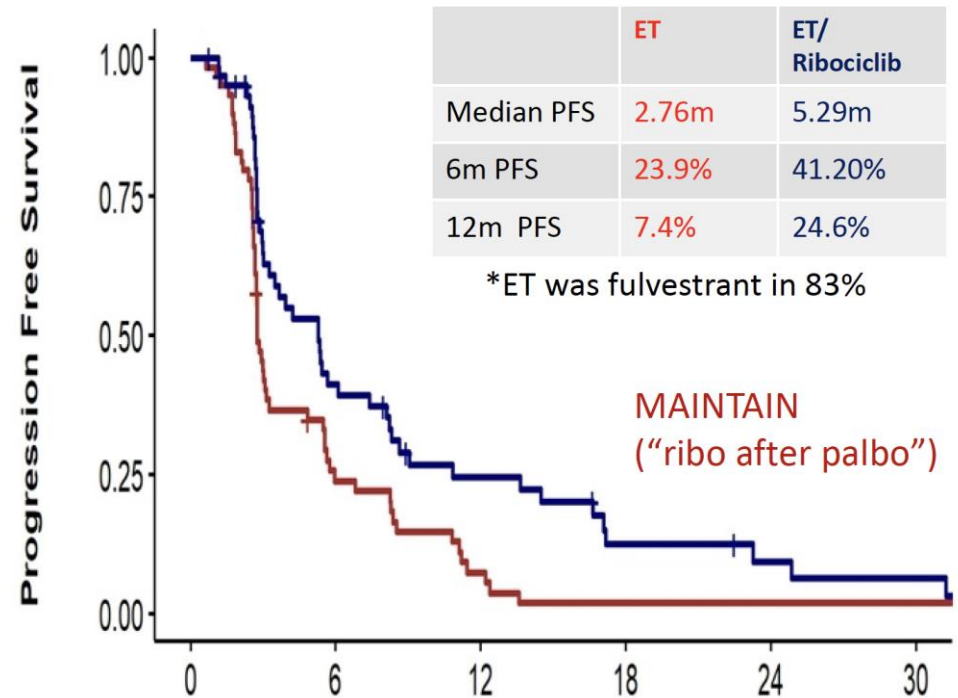
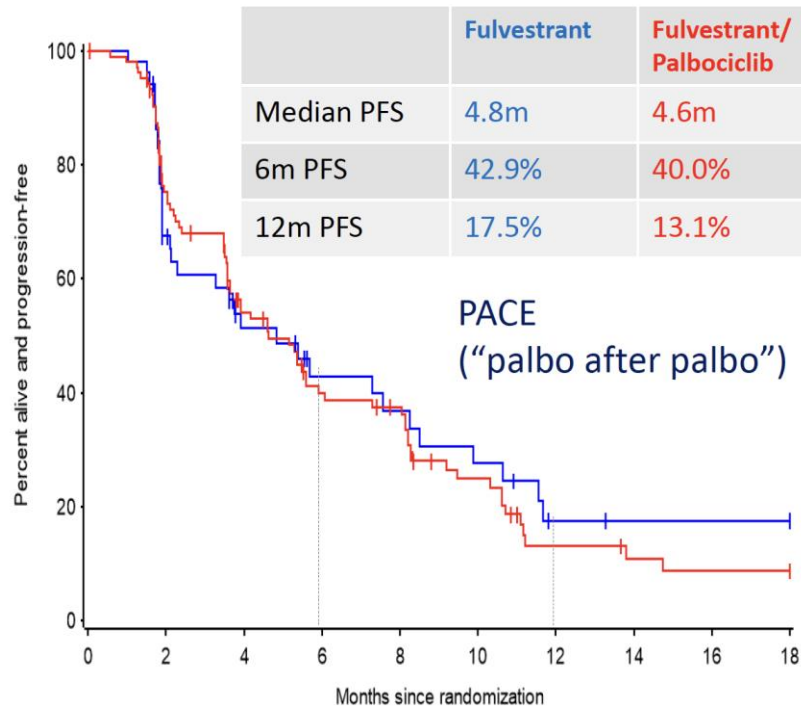
Basic cell-cycle regulation

Kim et al. 2023; Zhang et al. 2023;

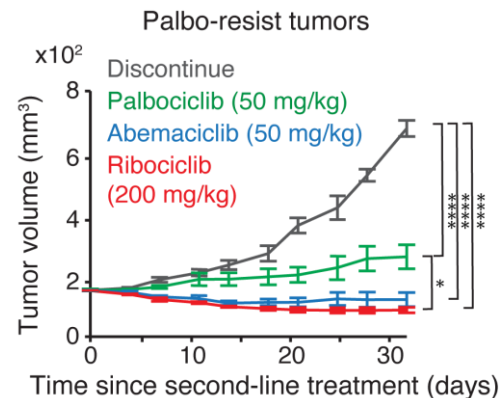
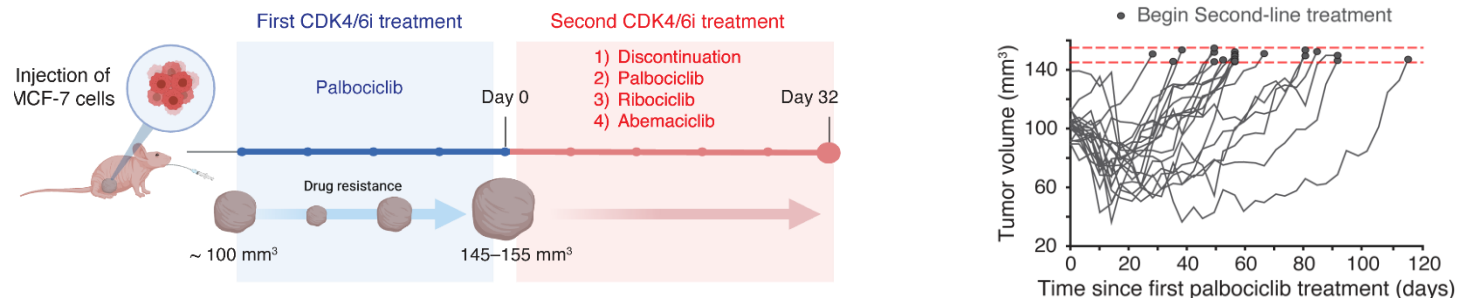
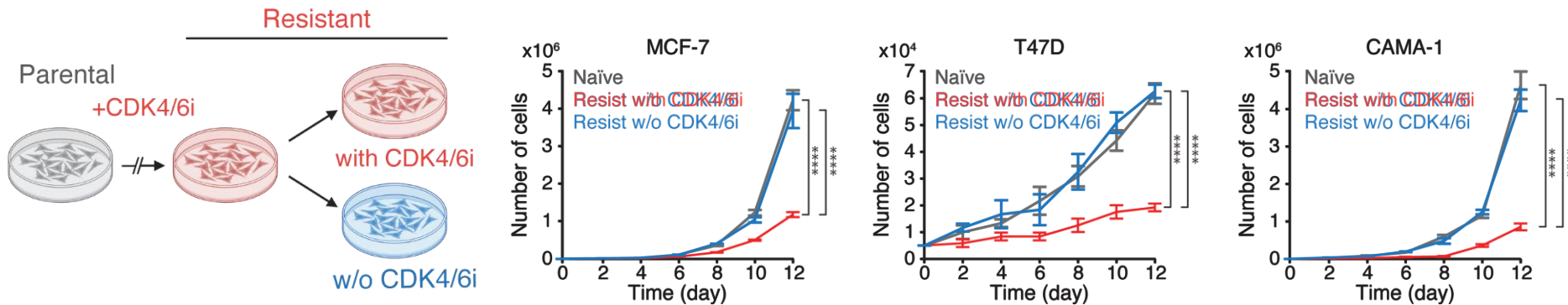


Continuing CDK4/6i beyond progression?

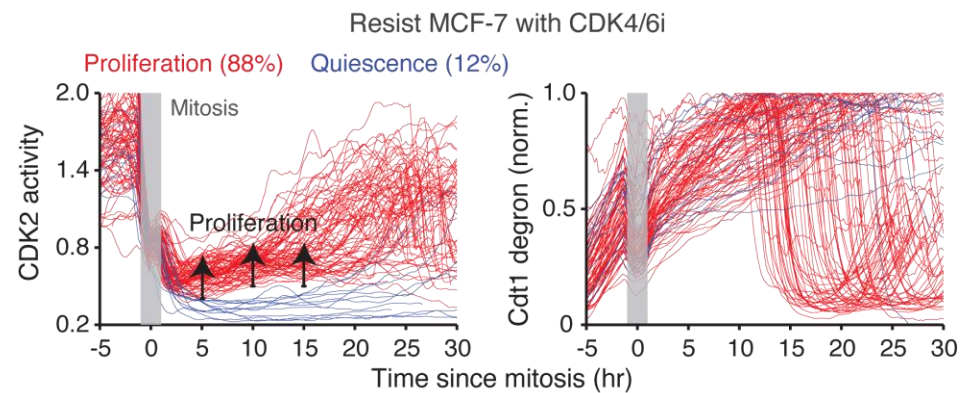
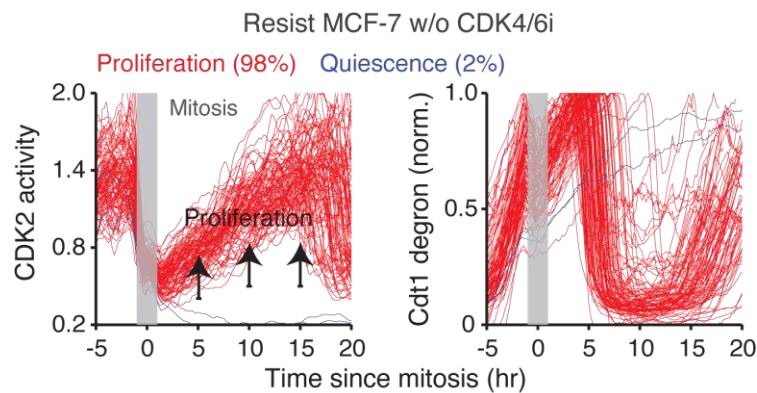
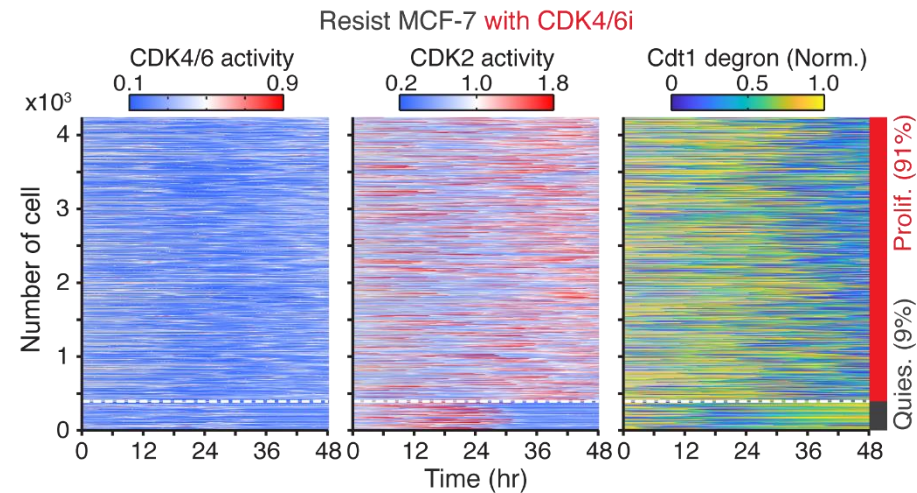
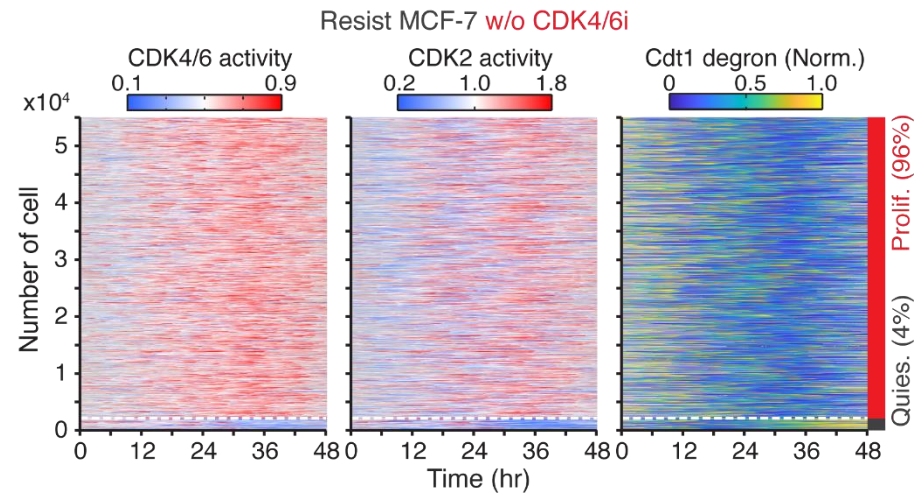
PACE and MAINTAIN clinical trials



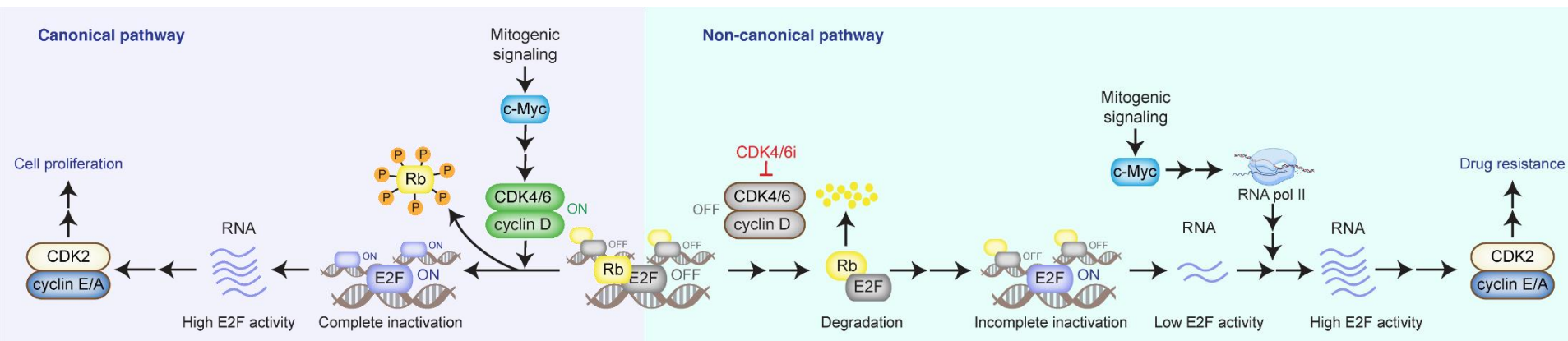
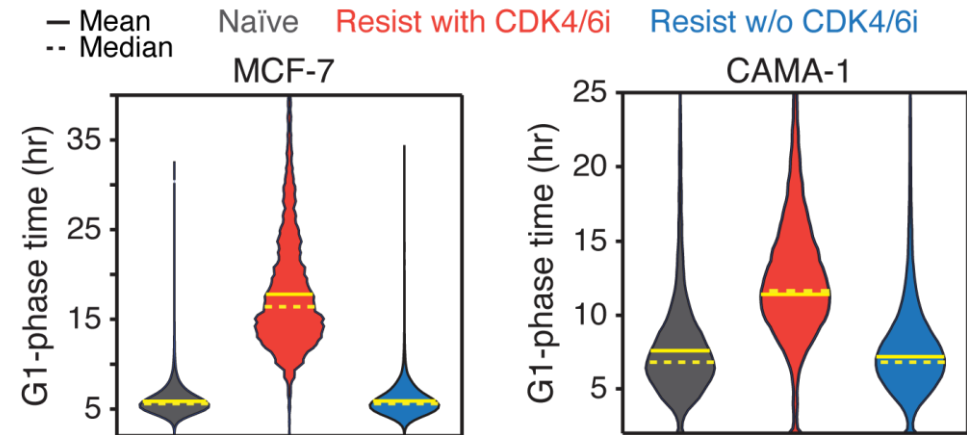
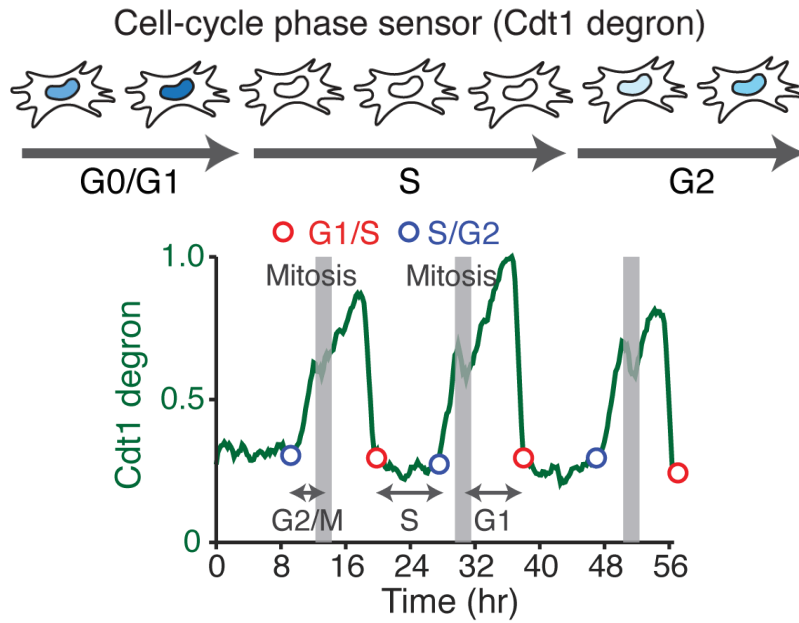
The benefits of CDK4/6i maintenance beyond disease progression



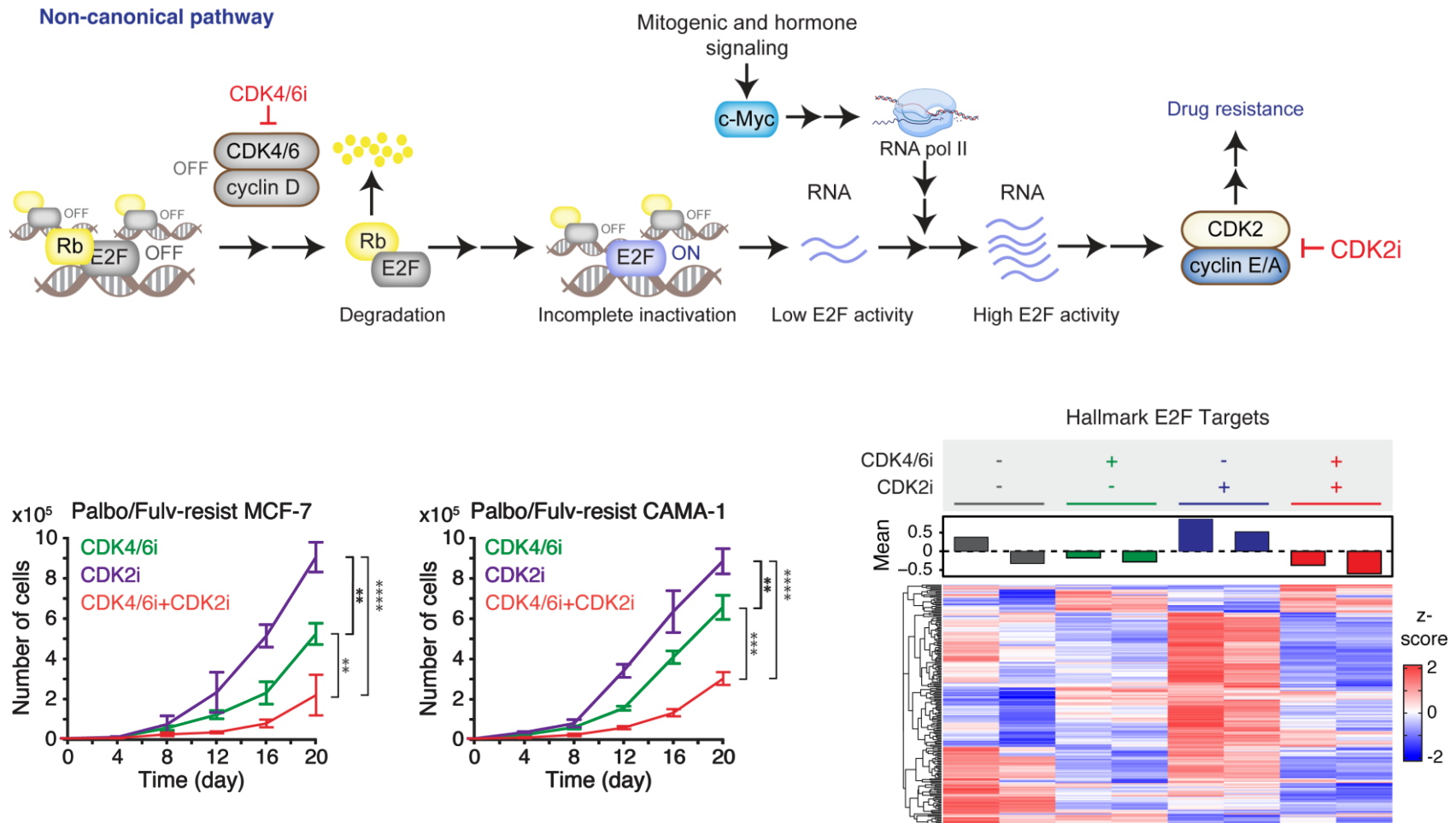
The benefits of CDK4/6i maintenance beyond disease progression



The benefits of CDK4/6i maintenance beyond disease progression



The benefits of CDK4/6i maintenance beyond disease progression



The role of CDK4/6 beyond the restriction point

Article

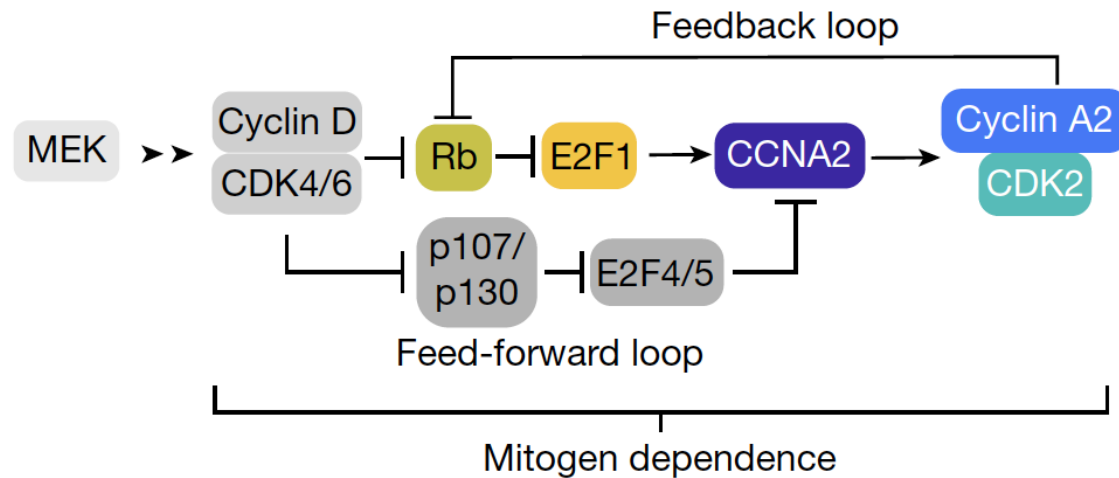
Loss of CDK4/6 activity in S/G2 phase leads to cell cycle reversal

<https://doi.org/10.1038/s41586-023-06274-3>

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James A. Cornwell¹, Adrijana Crncec¹, Marwa M. Afifi¹, Kristina Tang¹, Ruhul Amin¹ & Steven D. Cappel¹✉

Cornwell et al. 2023



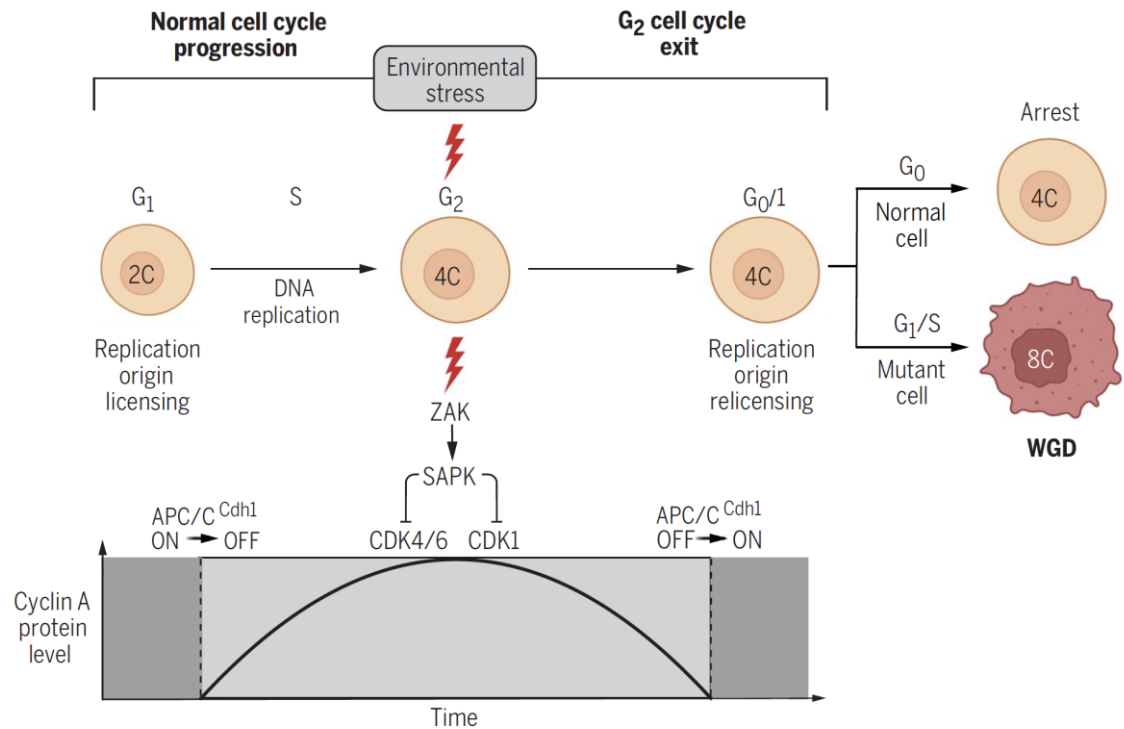
The role of CDK4/6 beyond the restriction point

CELL CYCLE

CDK4/6 activity is required during G₂ arrest to prevent stress-induced endoreplication

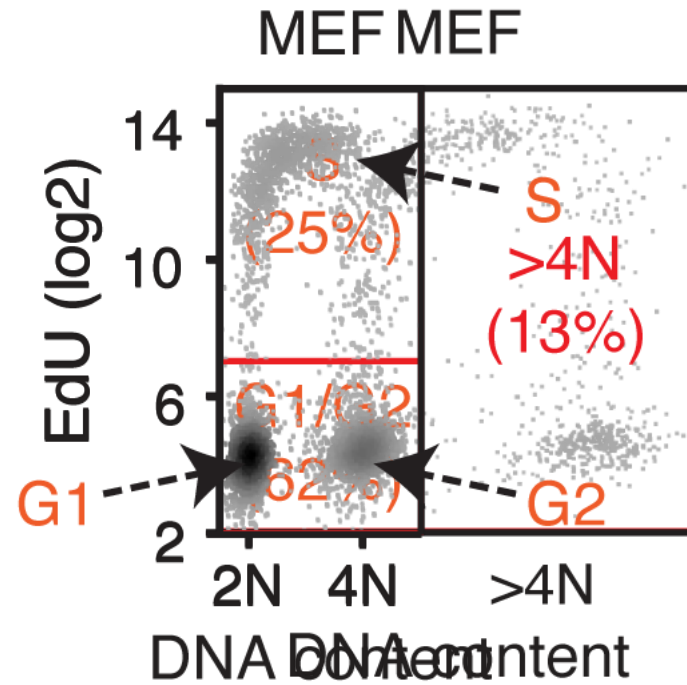
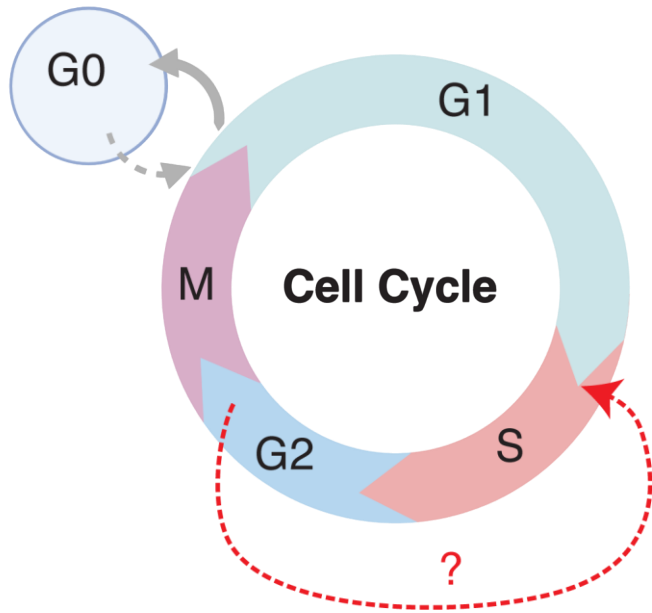
Connor McKenney, Yovel Lendner, Adler Guerrero Zuniga, Niladri Sinha, Benjamin Veresko, Timothy J. Aikin, Sergi Regot*

McKenney et al. 2024

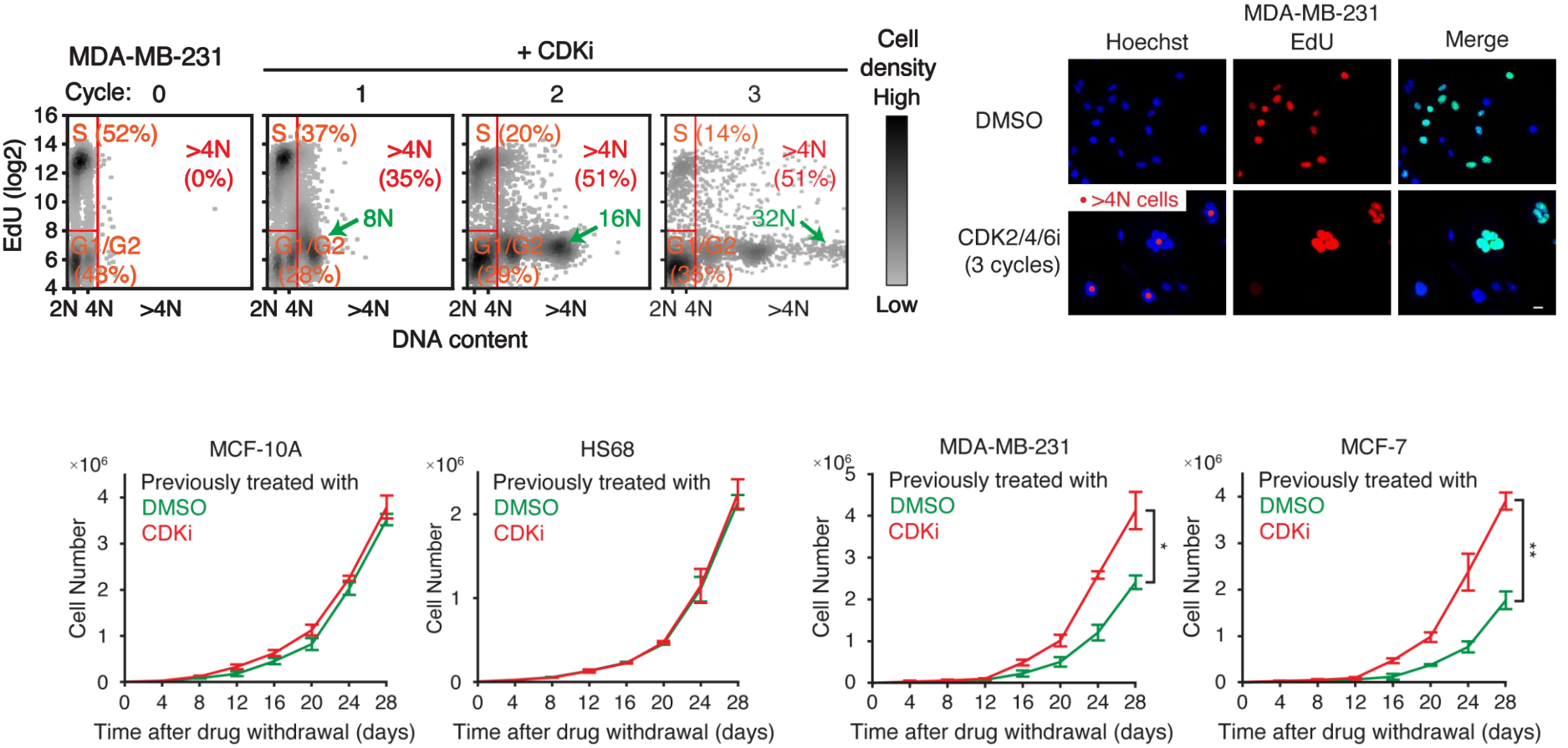


Whole-genome duplication

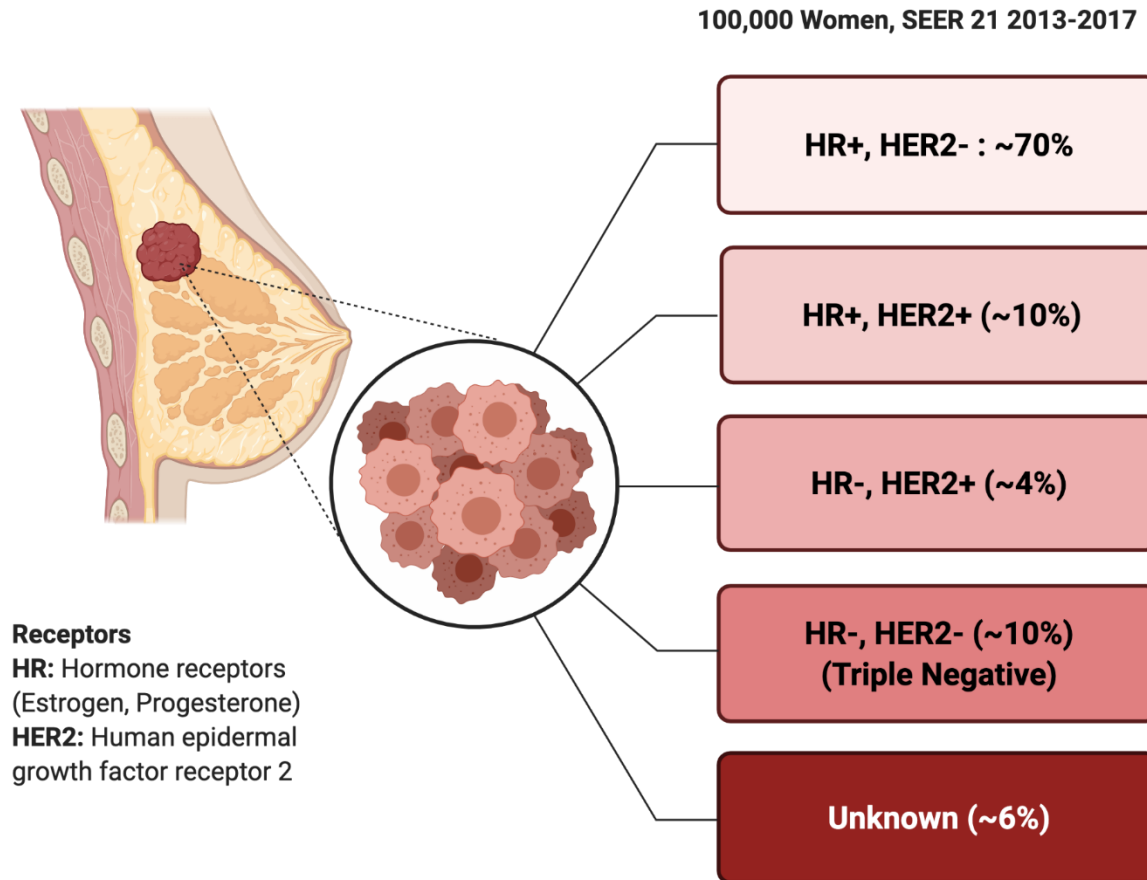
Kim et al. 2025



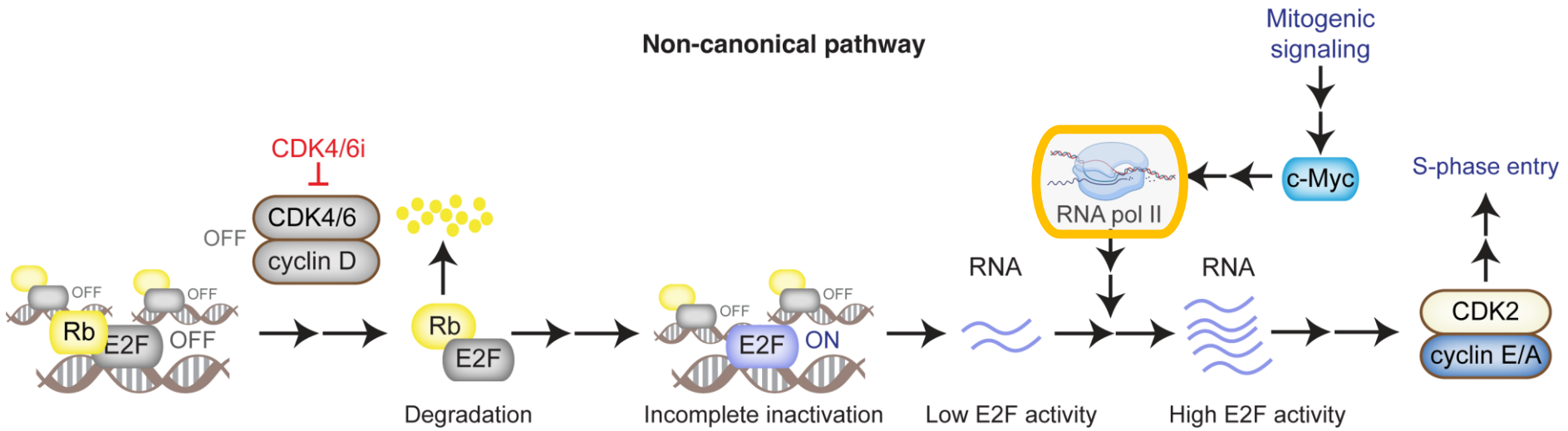
Consequences of WGD



Targeted therapeutic strategies for TNBC

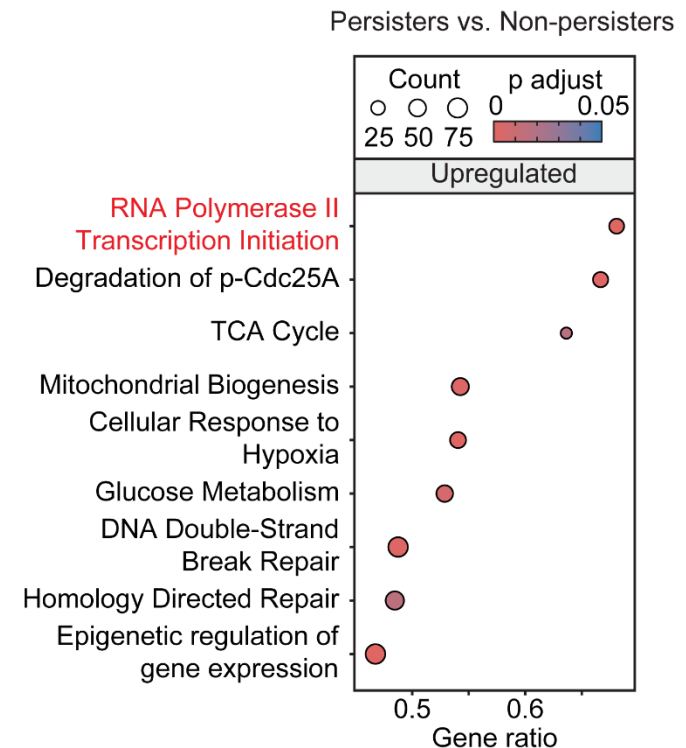
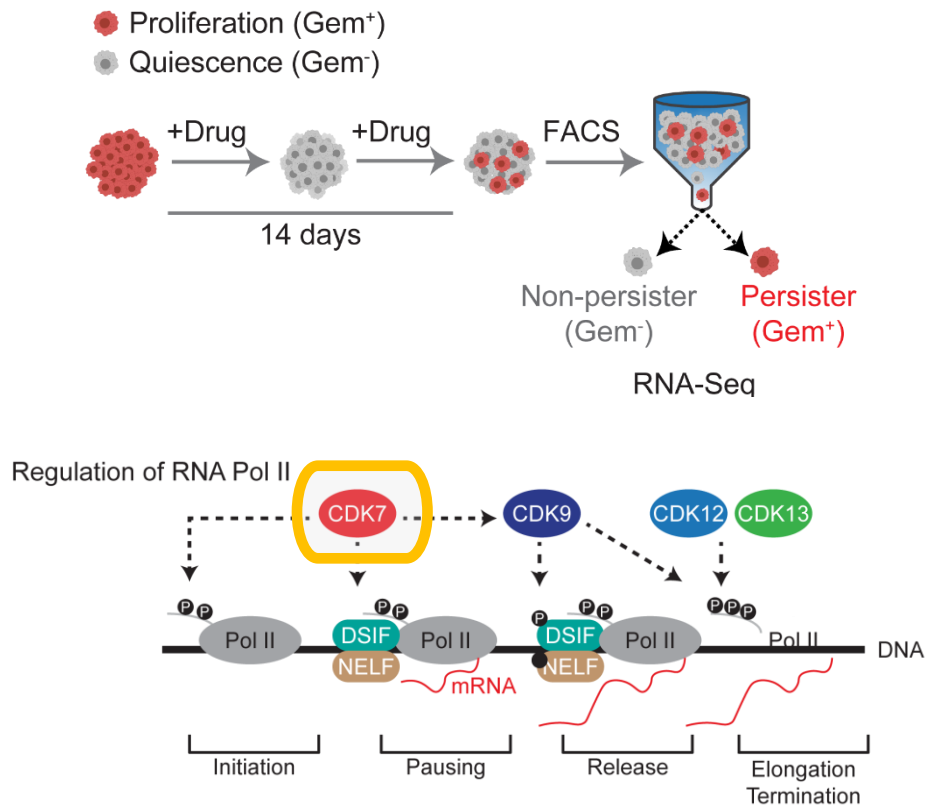


Targeting RNA polymerase II activity

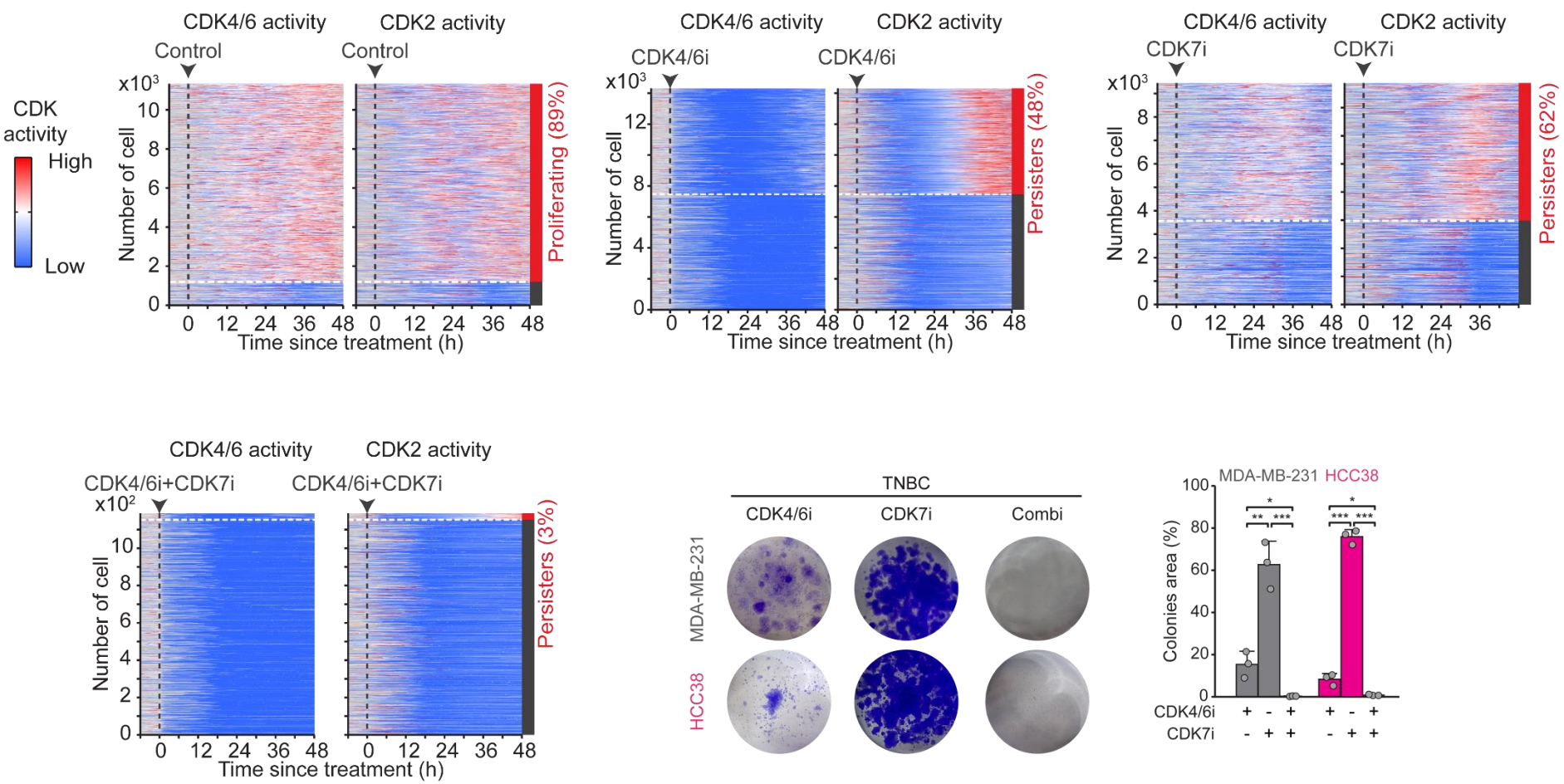


Lin et al., 2012; Nie et al., 2012; Rahl et al., 2010;

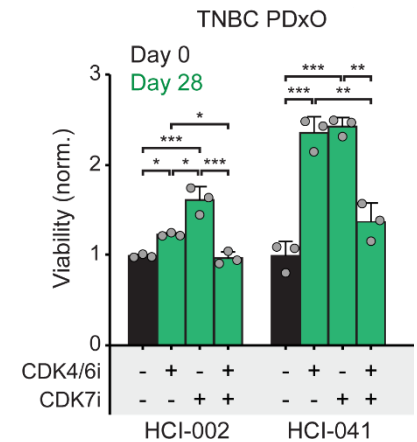
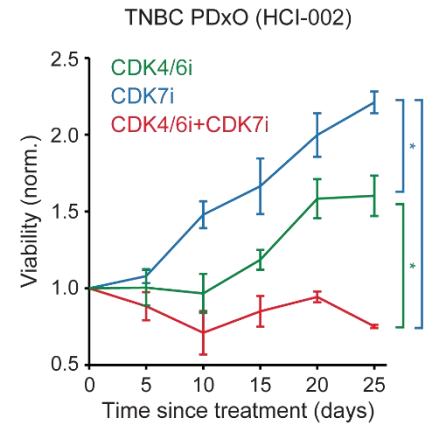
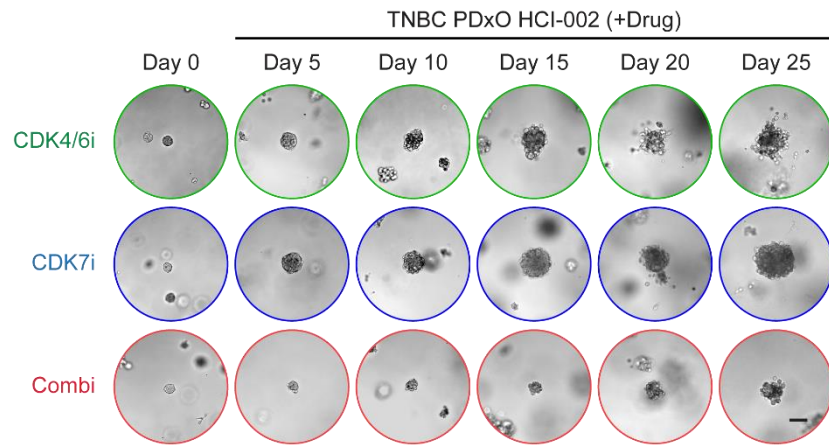
Targeting RNA polymerase II activity by CDK7 inhibitors



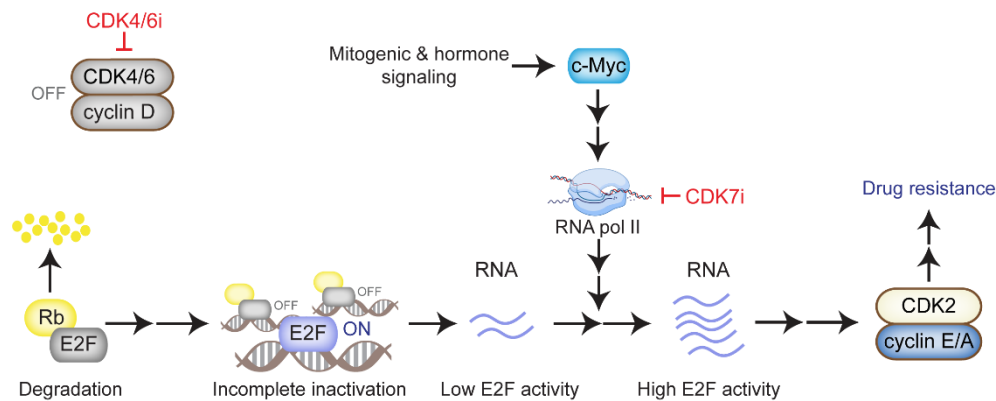
Dual targeting CDK4/6 and CDK7 activity (TNBC cell lines)



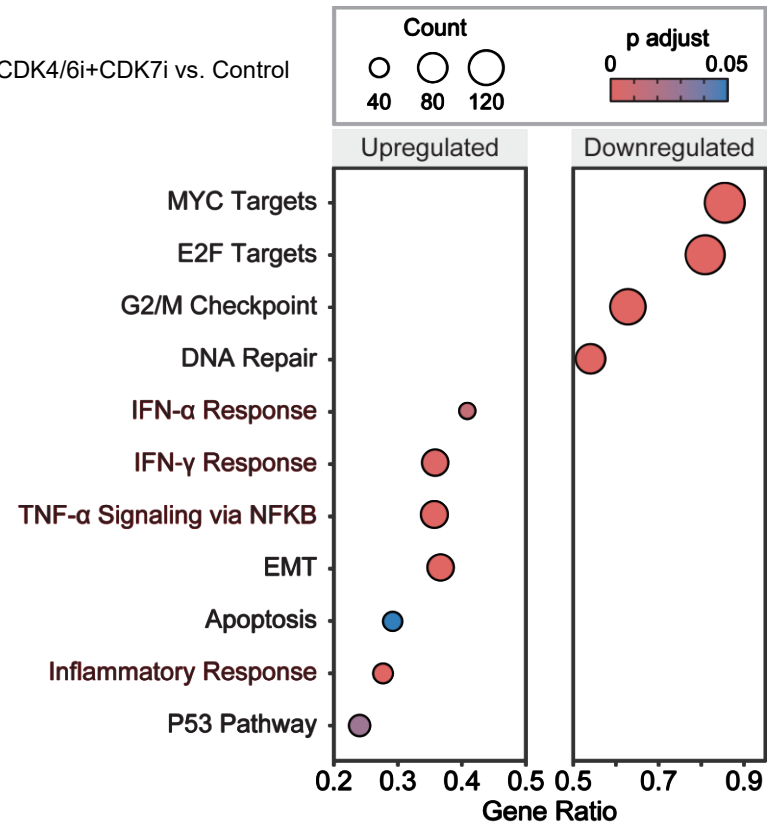
Dual targeting CDK4/6 and CDK7 activity (patient-derived organoids)



Upregulation of immune-related pathways

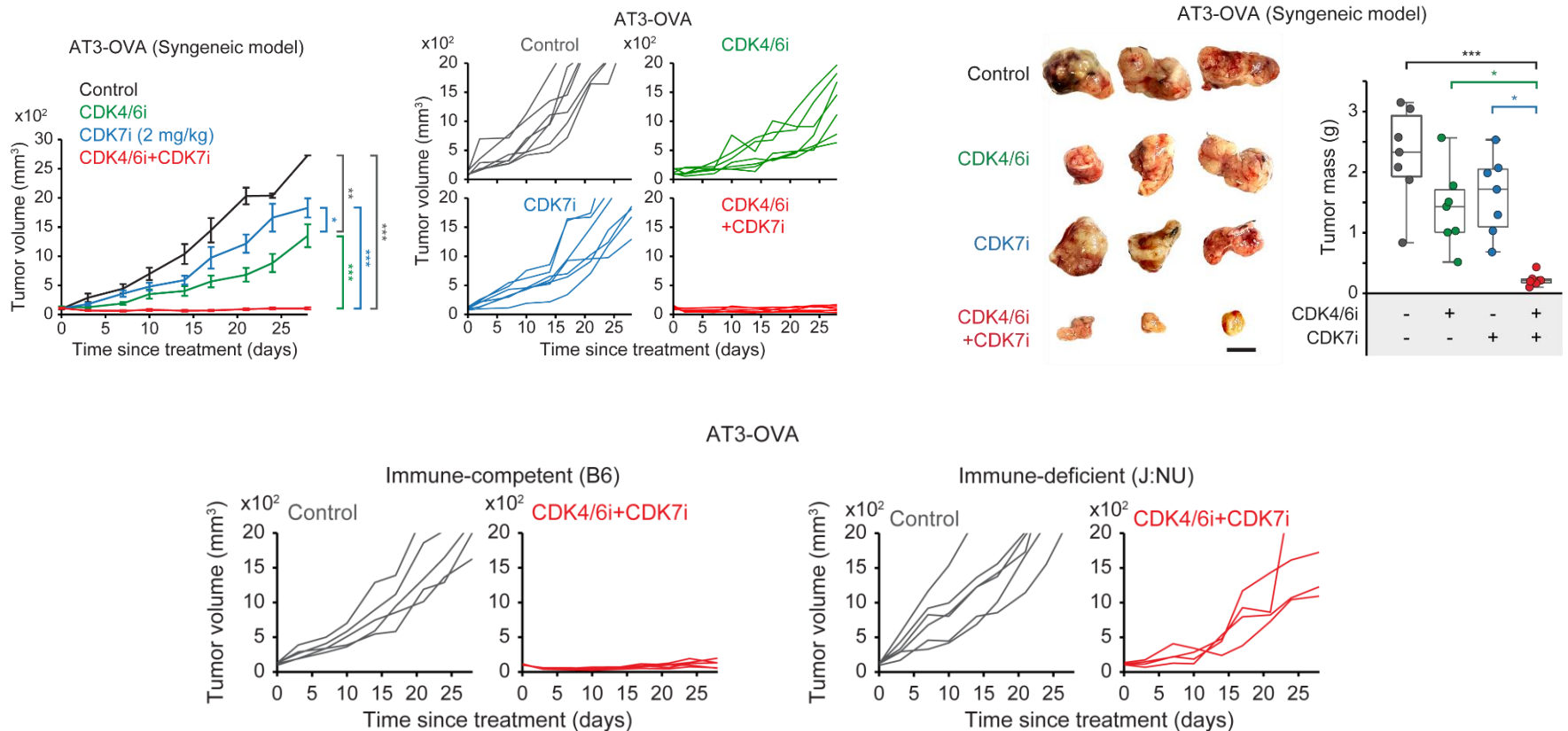


CDK4/6i+CDK7i vs. Control

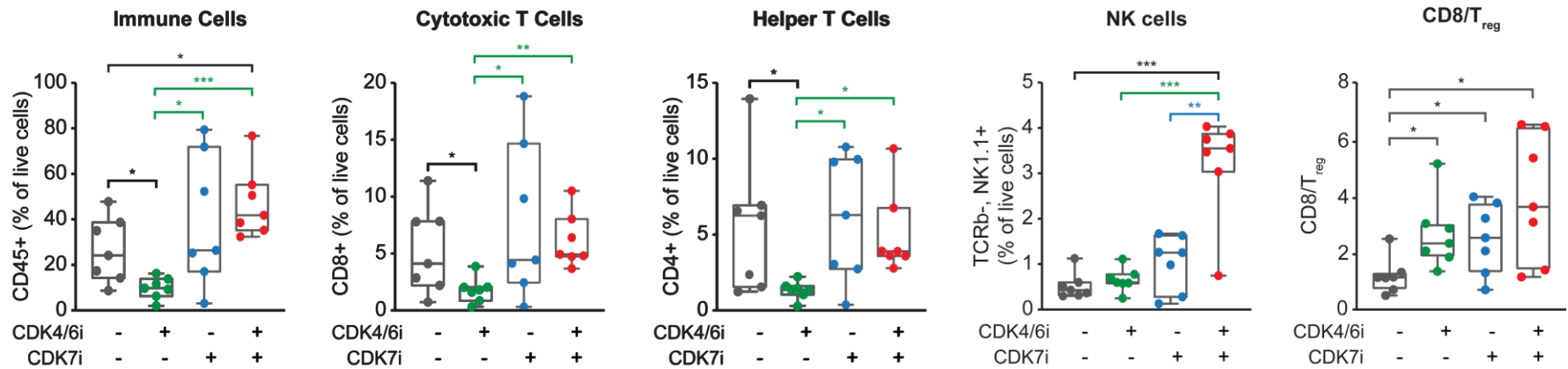


Dual targeting CDK4/6 and CDK7 activity (in vivo models)

$P < 0.01$, ** $P < 0.001$, *** $P < 0.0001$

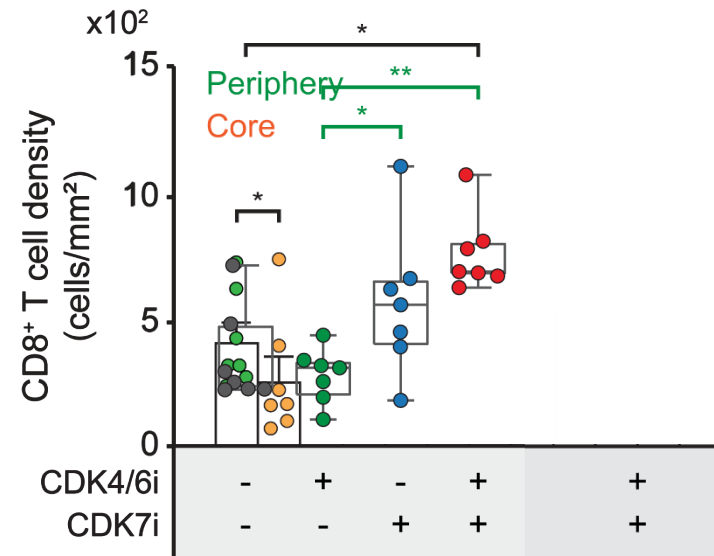
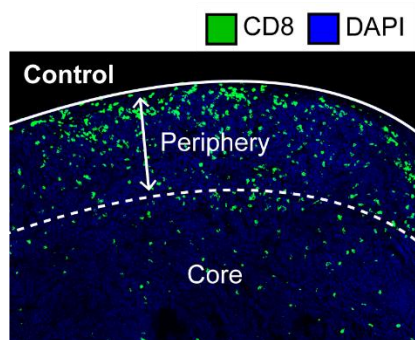


Impact on the tumor microenvironment

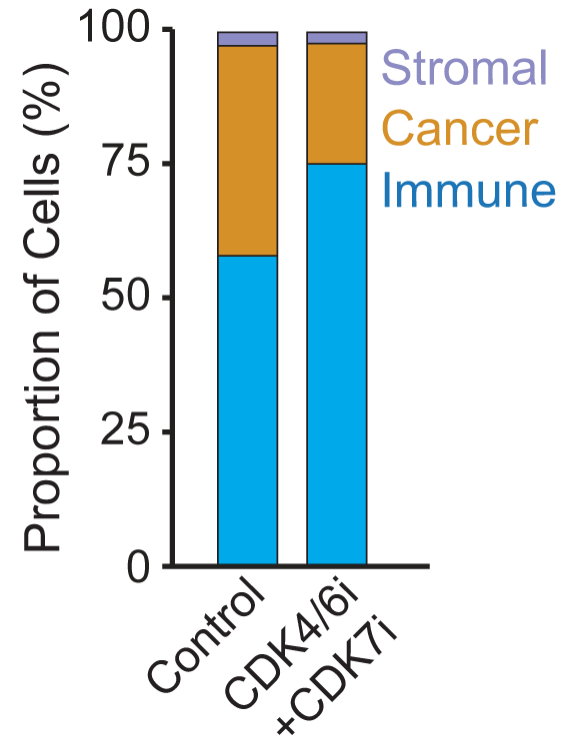
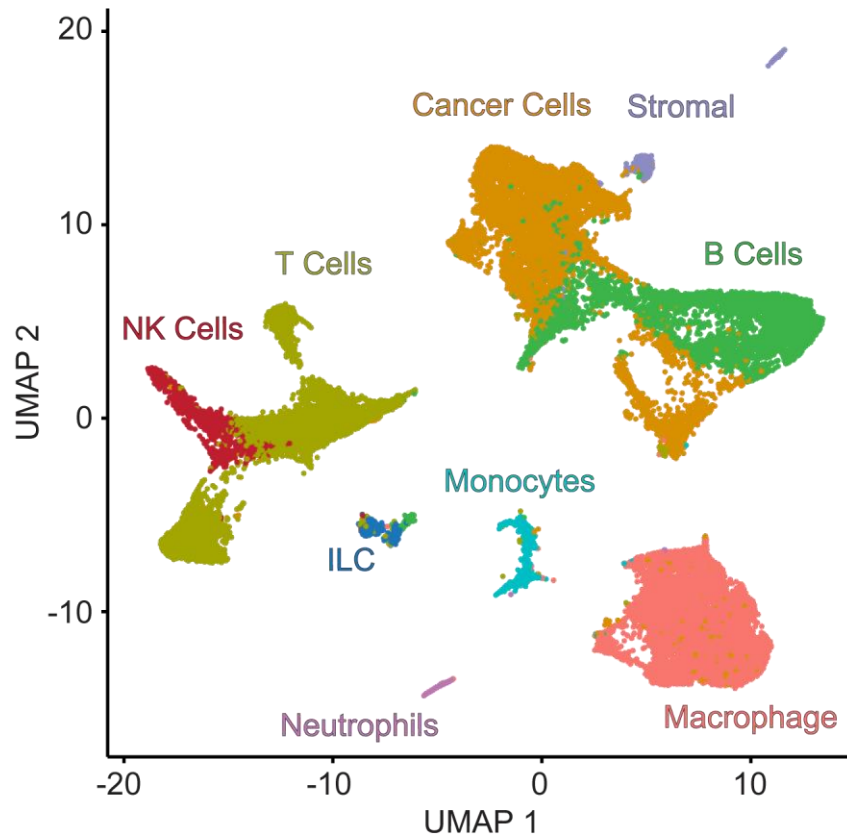


* P < 0.01, ** P < 0.001, *** P < 0.0001: one-way ANOVA

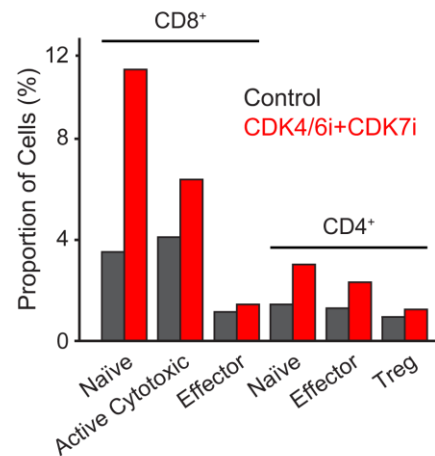
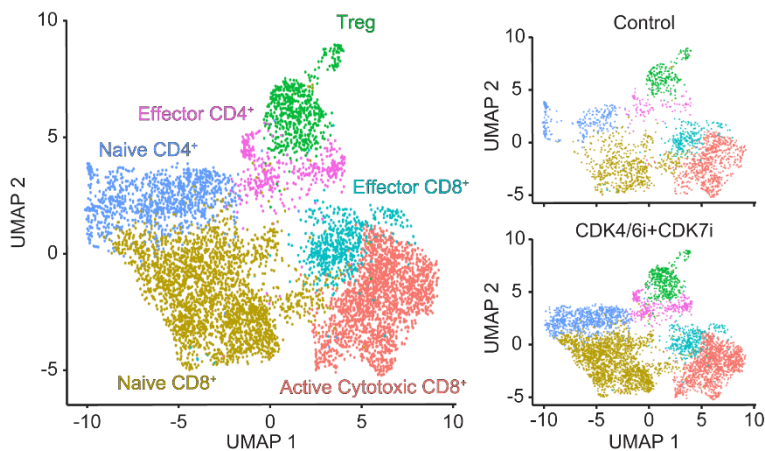
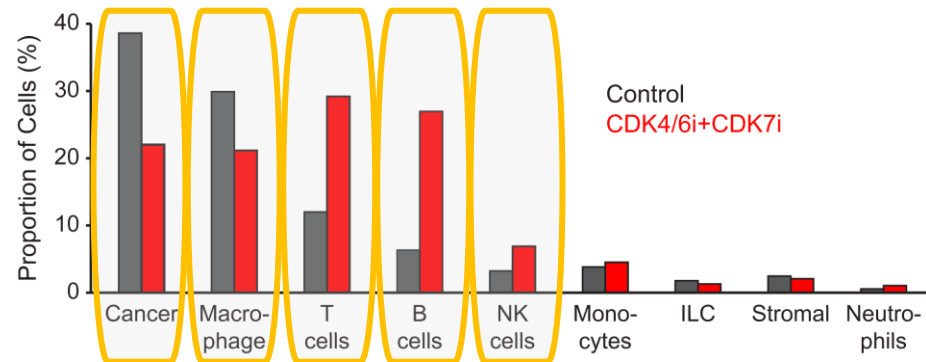
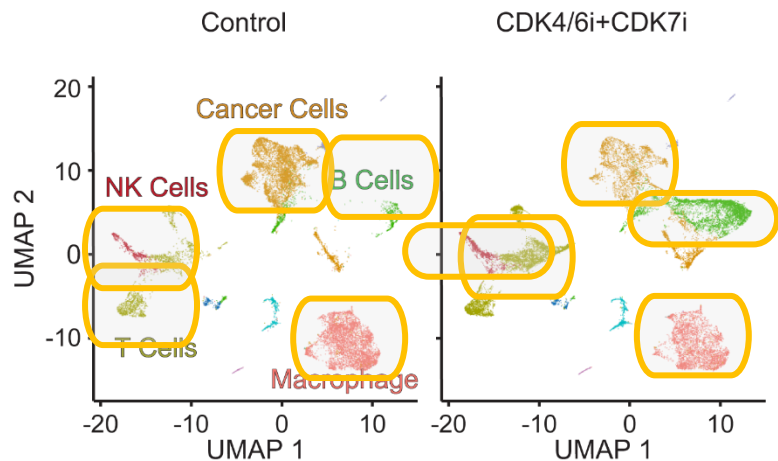
Impact on the tumor microenvironment



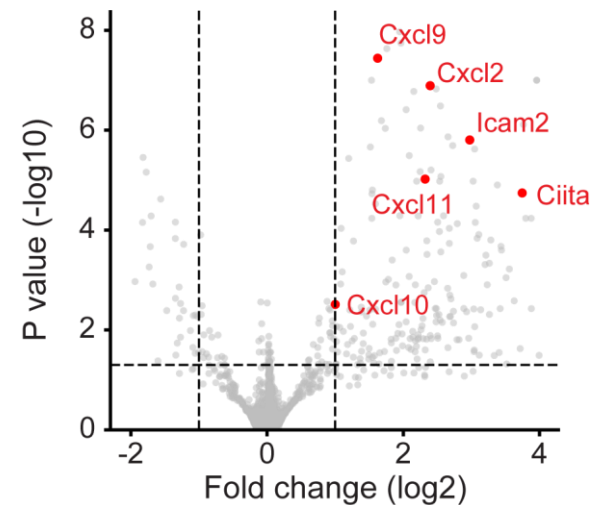
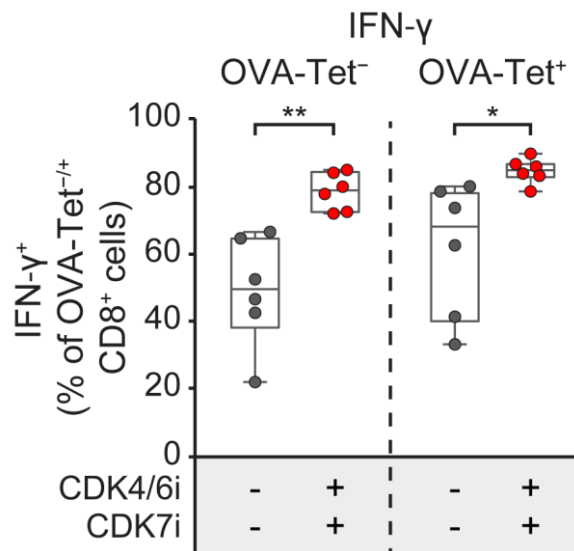
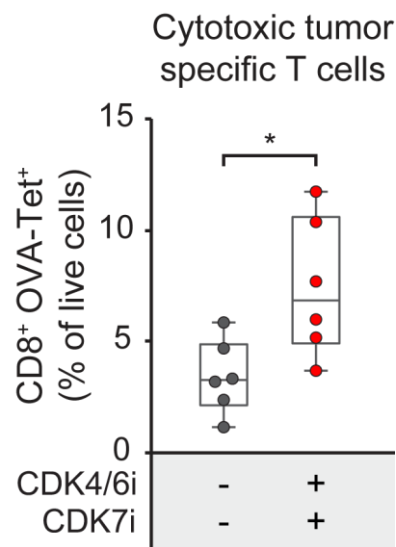
Impact on the tumor microenvironment



Impact on the tumor microenvironment

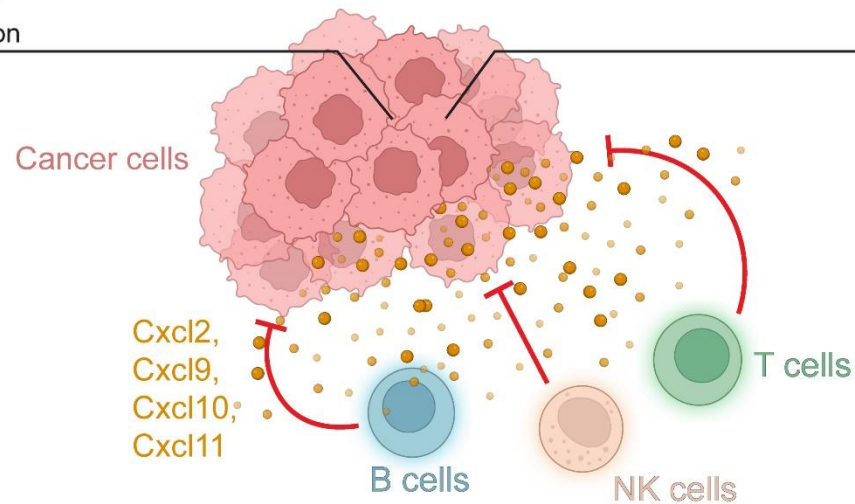
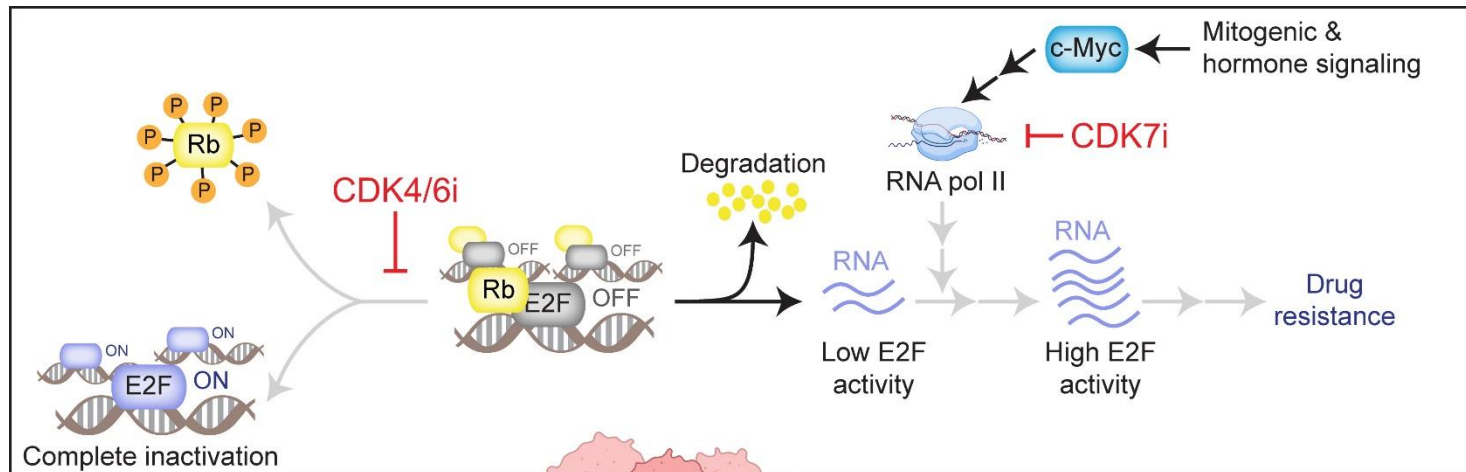


Impact on the tumor microenvironment

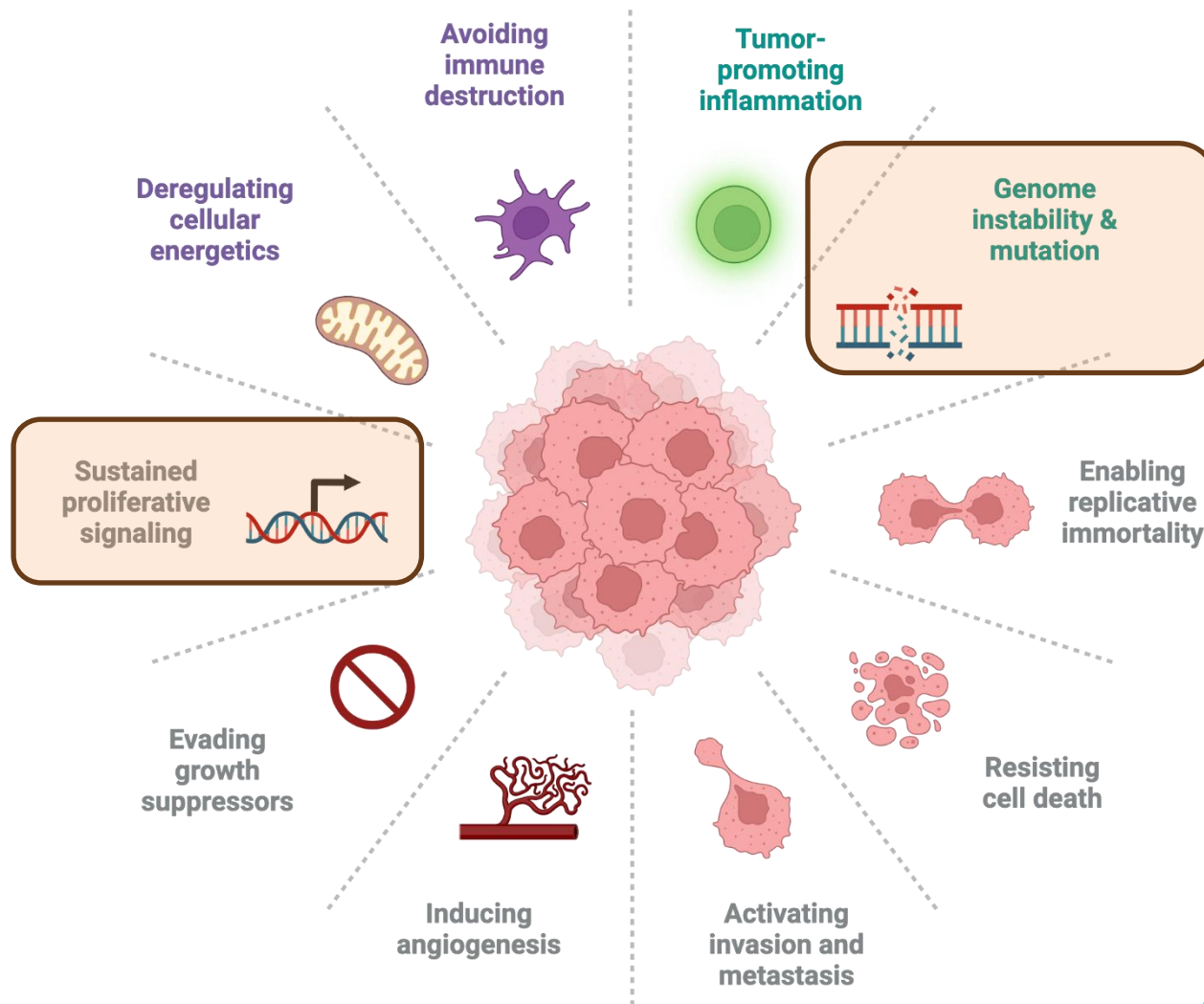


New Therapeutic strategy based on basic cell-cycle studies

Kim* and Son* et al., 2025

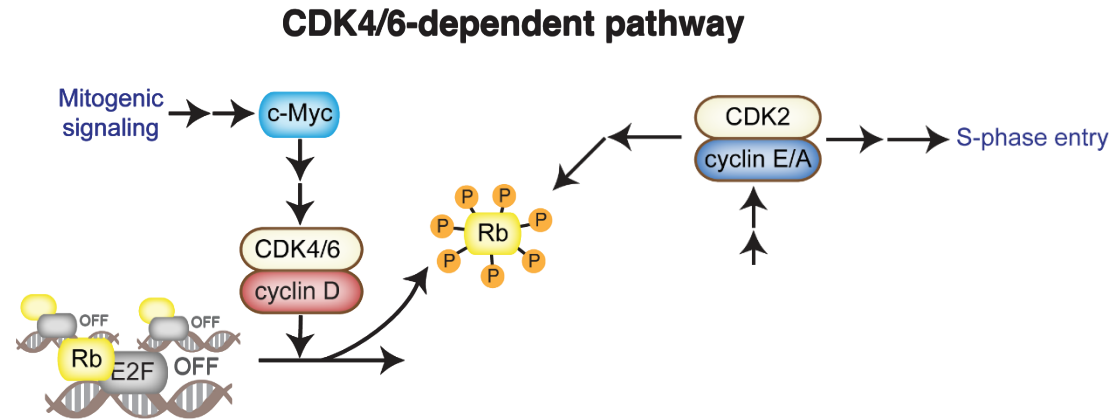
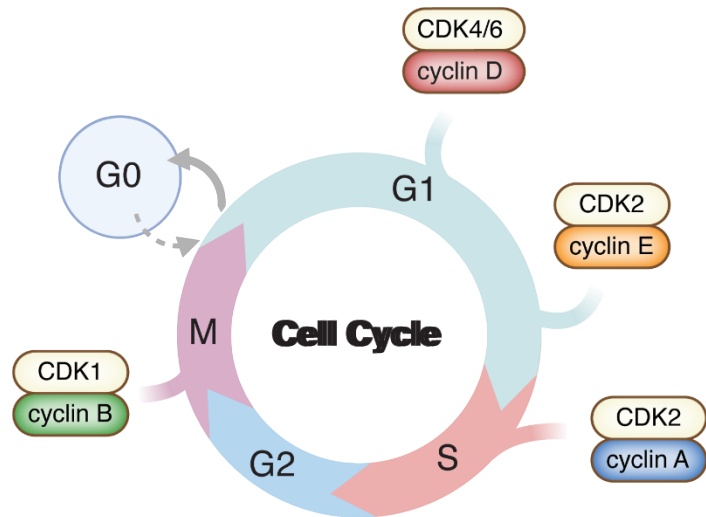


Hallmarks of cancer



Original hallmarks
Enabling factors
Emerging hallmarks

Cell cycle regulation



Any questions?

