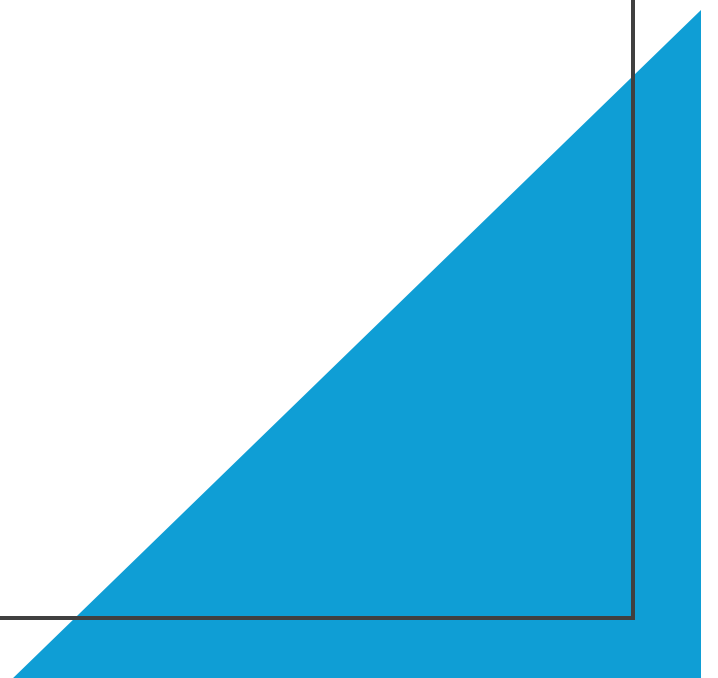


Preclinical Models of Cancer

PATH4500: Cancer Biology I

October 27, 2025

Carla Concepcion-Crisol, PhD



Goal: To bridge basic discovery and clinical application

Challenge: How do we model the complexity of human cancer?

“All models are wrong, but some models are useful.”

George E.P. Box

Key Learning Objectives



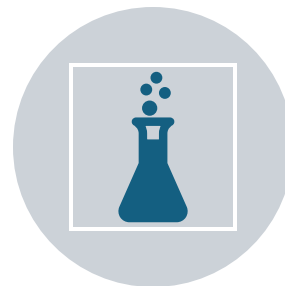
Understand various preclinical models of cancer



Assess the strengths and weaknesses of these models



Make informed decisions about which model is most suitable to use for what question



Design preclinical studies

Preclinical models of cancer

- *In vitro*
 - Cell lines
 - Spheroids
- *Ex vivo*
 - Organoids
 - Organ-on-a-chip
- *In vivo*
 - Genetically engineered mouse models (GEMMs)
 - Patient-derived xenografts (PDXs)
 - Syngeneic and transplant models
- (Deeper) dive into lung cancer models
- Research-in-progress examples

Some considerations

Fidelity to the biology

- Tumor initiation, progression, invasion, metastasis
- Therapeutic response
- Histopathology
- Tumor-stroma interactions
- Immunocompetence
- Secondary mutations
- How faithful to human biology?

What question are you asking?

Some considerations

Practical

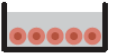
- Speed
- Scale
- Cost
- Ease of genetic manipulation

Is the model “good enough”?

Fidelity and practicality often compete

Table 1. Properties of cancer model systems.

GEMM, genetically engineered mouse model; MDO, murine-derived organoid; MDOX, murine-derived organoid transplantation; CLs, cell lines; PDX, patient-derived xenograft; iPS, inducible pluripotent stem cell; PDO, patient-derived organoid; PDOX, patient-derived organoid transplantation.

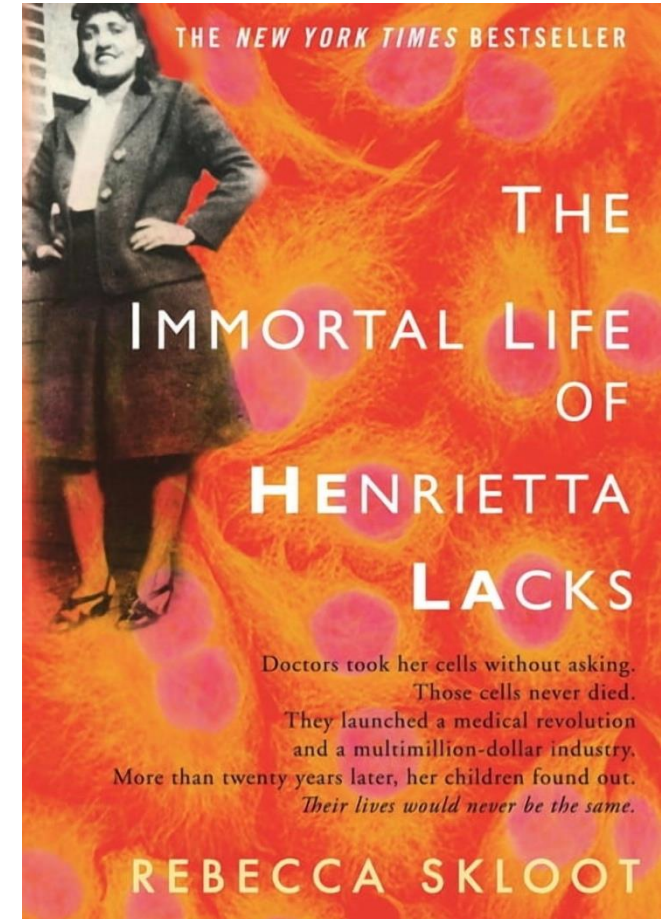
	 GEMM	 MDO	 MDOX	 CLs	 PDX	 iPS	 PDO	 PDOX
Wild-type cell culture	+	+	+	–	–	+	+	–
Preinvasive cancer models	+	+	+	–	–	+	+	+
Invasive cancer models	+	+	+	+	+	+	+	+
Metastatic cancer models	+	+	+	+	+	+	+	+
Cost	\$\$\$\$	\$\$	\$\$\$	\$	\$\$	\$\$	\$\$	\$\$\$
Time	++++	+	++	+	++++	+++	++	+++
Success rate	high	med	med	med	med	low	med	med
Throughput therapies	low	med	low	high	low	high	med	low

+ denotes 1 month or less; ++, 1–2 months; +++, 1–6 months; +++++, oftentimes more than several months.

Cell lines

1951: The first human cancer cell line, HeLa, was established from cervical cancer cells of Henrietta Lacks without her knowledge or consent

1990s: The National Cancer Institute launched the NCI-60 project, which led to a panel of 60 human cancer cell lines of various types



Cell lines



Towards mapping the landscape of cancer vulnerabilities across all tumors

[Explore the DepMap public data →](#)

[DMC Access](#)

<https://depmap.org/portal/home/#/>
<https://sites.broadinstitute.org/ccle/>
<https://www.cancerrxgene.org>



Genomics of Drug Sensitivity in Cancer

Home Compounds Genes Cell Lines About News Downloads Documentation

The Genomics of Drug Sensitivity in Cancer project is an academic research program to identify molecular features of cancers that predict response to anti-cancer drugs.

Browse our data

 [Compounds](#)

 [Cancer Genes](#)

 [Cell Lines](#)

Please be aware that our website and results are part of an ongoing project. These webpages are updated frequently and our results are not final or complete.

Search

Enter drug, gene or cell line

e.g. Docetaxel, RP-56976, BRAF, COLO-829

What's new!

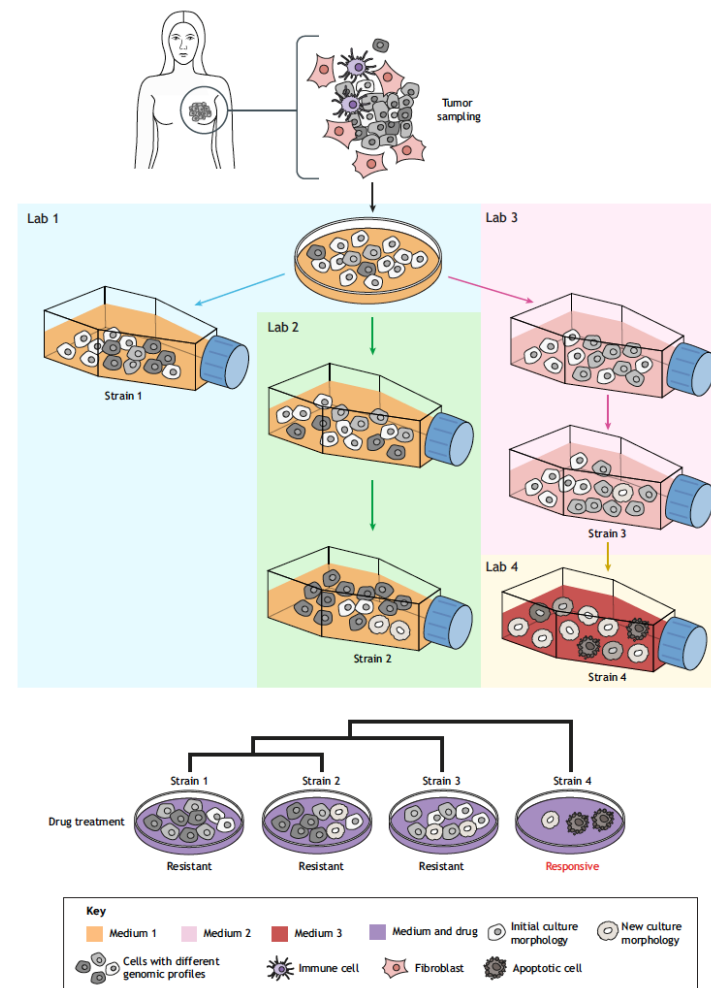
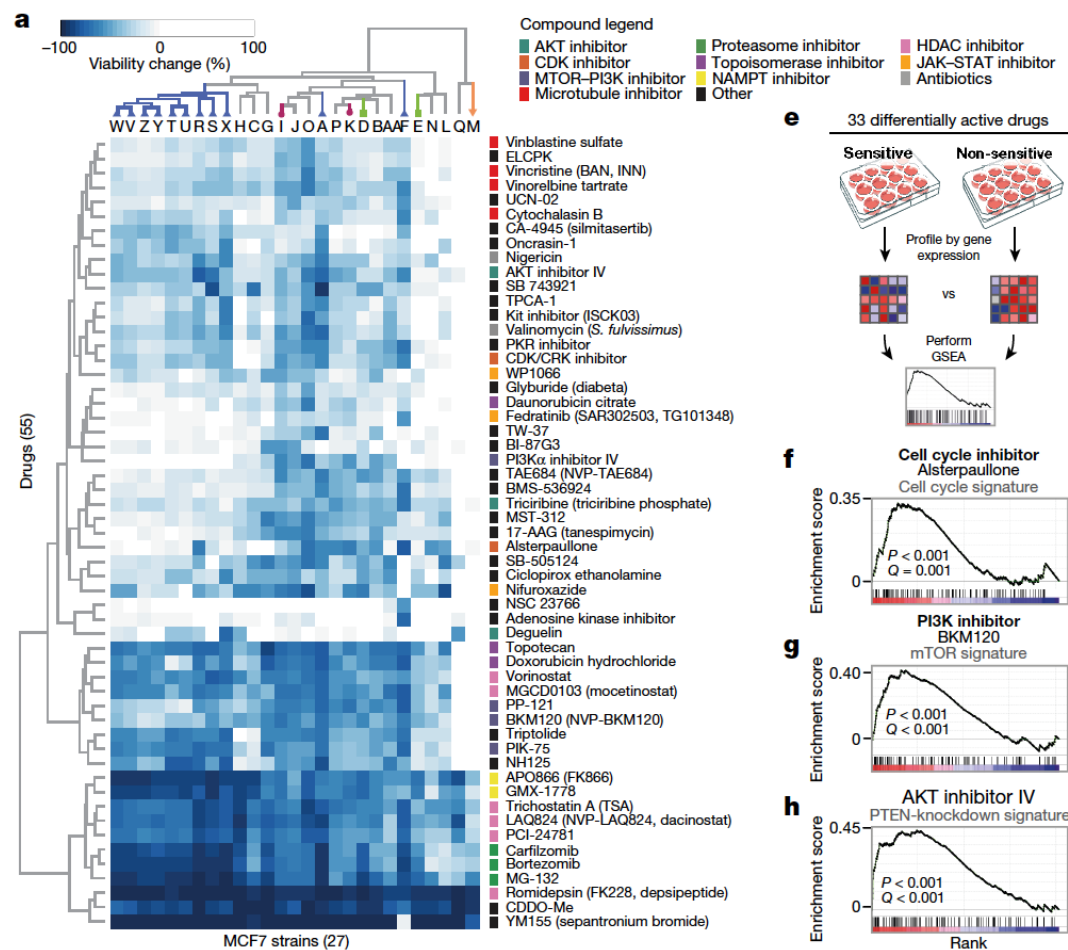
Release 5 (June 2014)

Sensitivity data can now be browsed by cell line. The cell line page shows the sensitivity across all drugs for each cell line using a z-score value.

Mailing List

To receive news and data release alert, please sign up [Translation-announce](#)

Cell lines



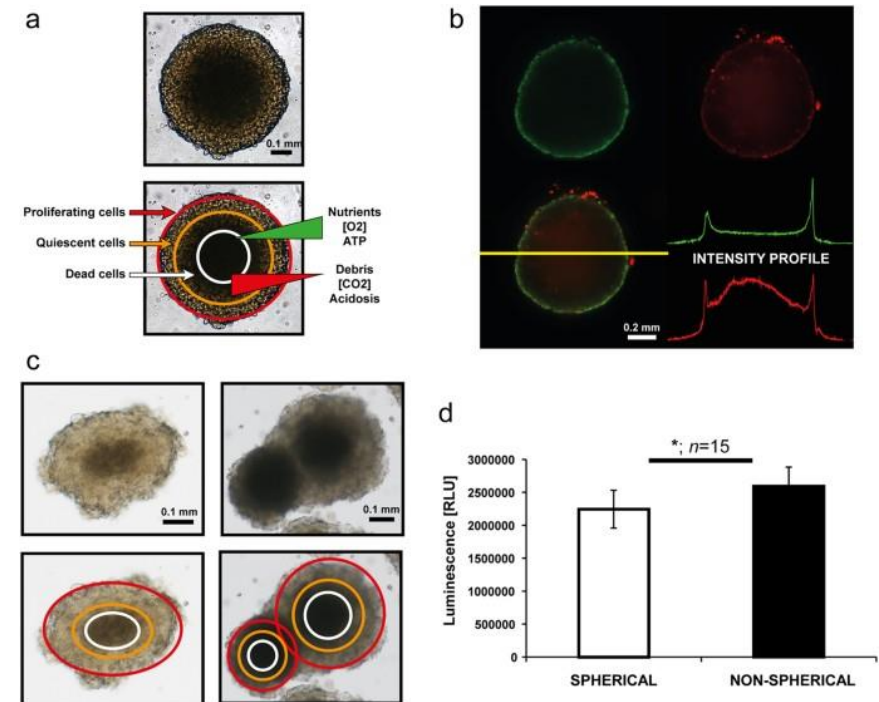
2D to 3D: cancer spheroids

3D aggregates of immortalized cancer cell lines

Often uses ultra-low adhesion dishes

Some form without addition of extracellular matrix (ECM) proteins; others require the addition of basement membrane extract (BME) or collagen for efficient formation

Mimic cell-cell interaction, hypoxia, drug penetration, response and resistance and production/deposition of extracellular matrix to an extent greater than 2D cultures

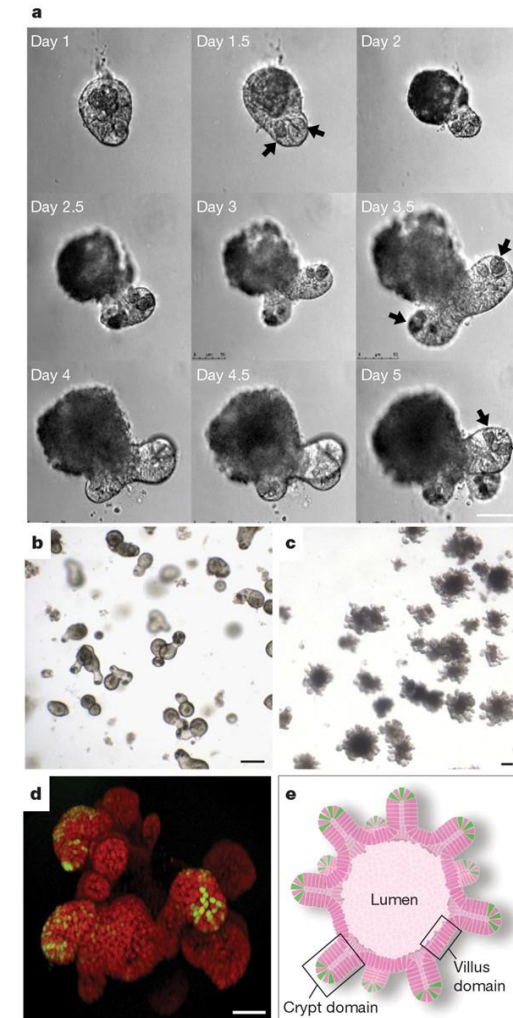


Organoids represent a breakthrough in modeling normal tissues

Microscopic self-organizing, 3D structures that are grown from stem cells *in vitro*

Recapitulate many structural and functional aspects of *in vivo* counterpart organs

Adapted to culture tumor organoids of various cancer types



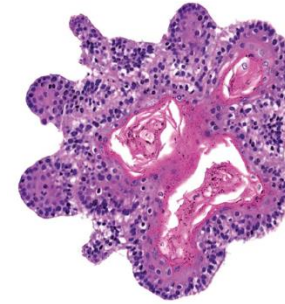
Patient-derived organoids (PDOs)

Mimic the 3D structure and function of human tissues

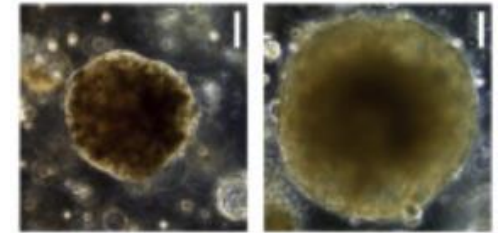
Capture the cellular heterogeneity and spatial organization found within tumors

Cultures with stromal cells enable studies of the tumor microenvironment (TME)

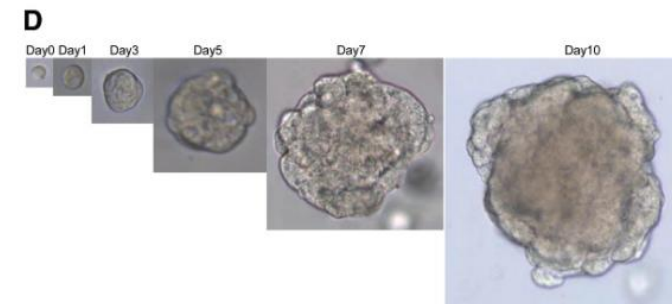
Uses extracellular matrices, air-liquid interface (ALI) cultures, chips, microfluidics



Squamous cell carcinoma (oral mucosa)

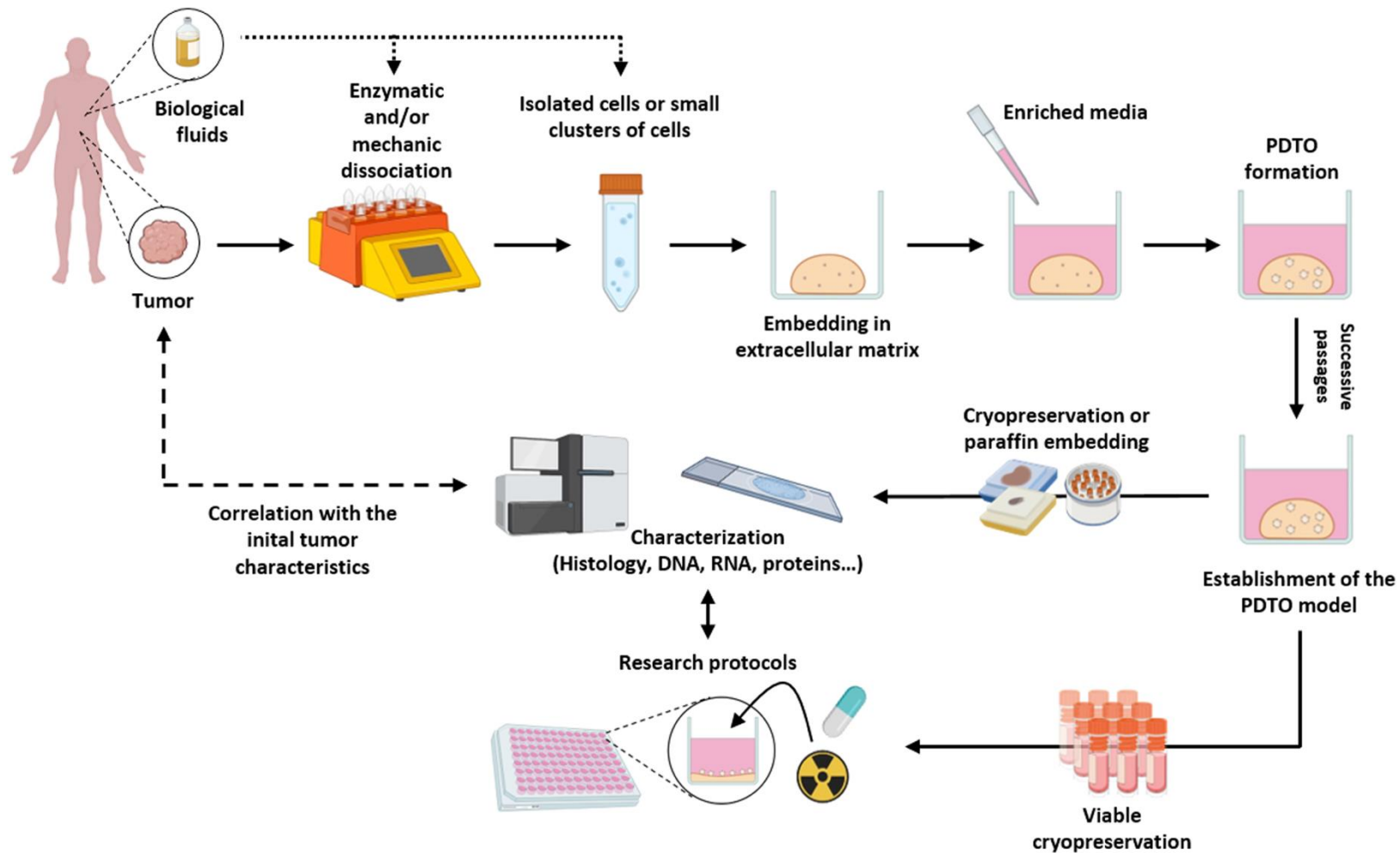


Bladder cancer

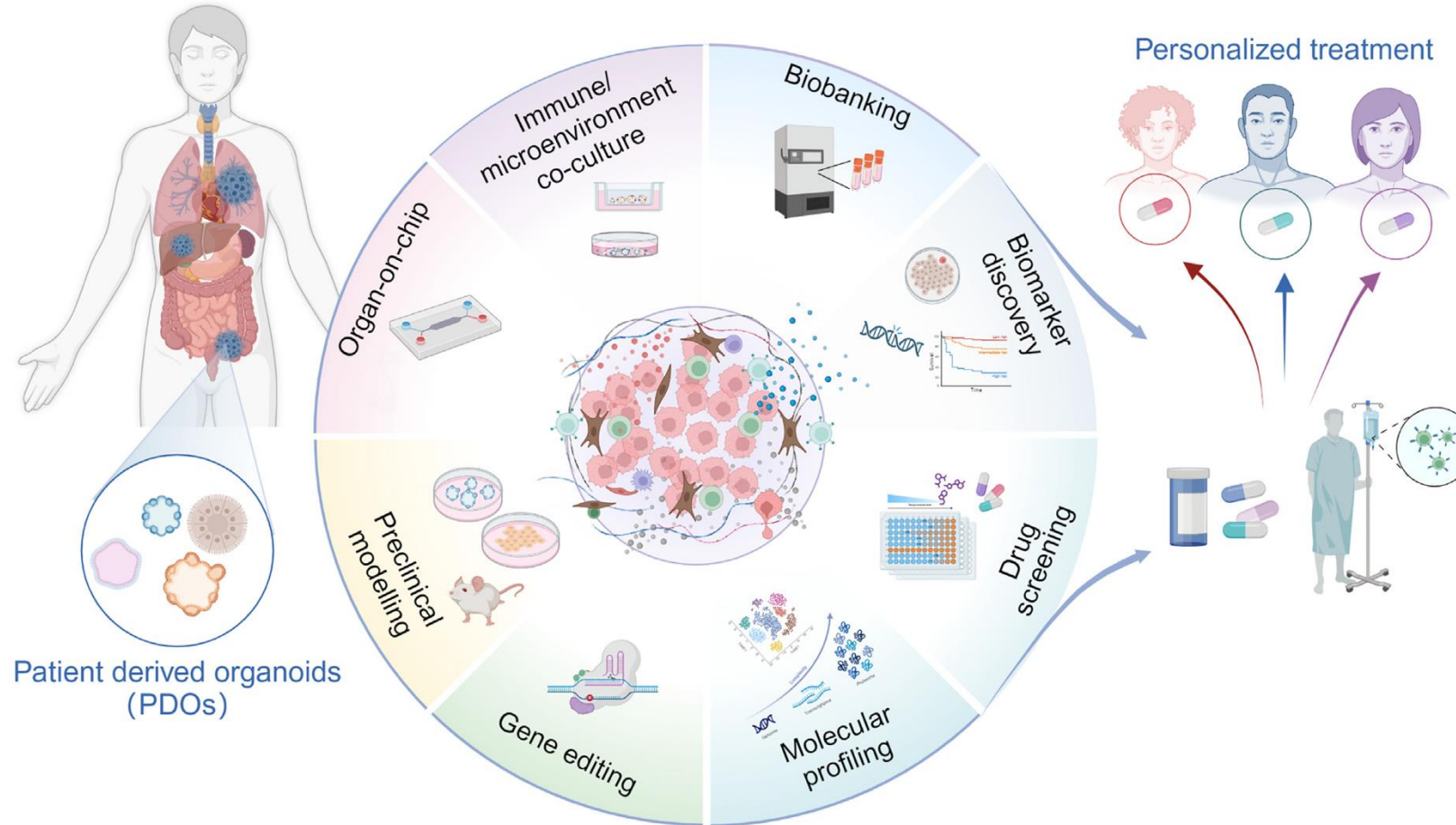


Human colon adenocarcinoma organoids

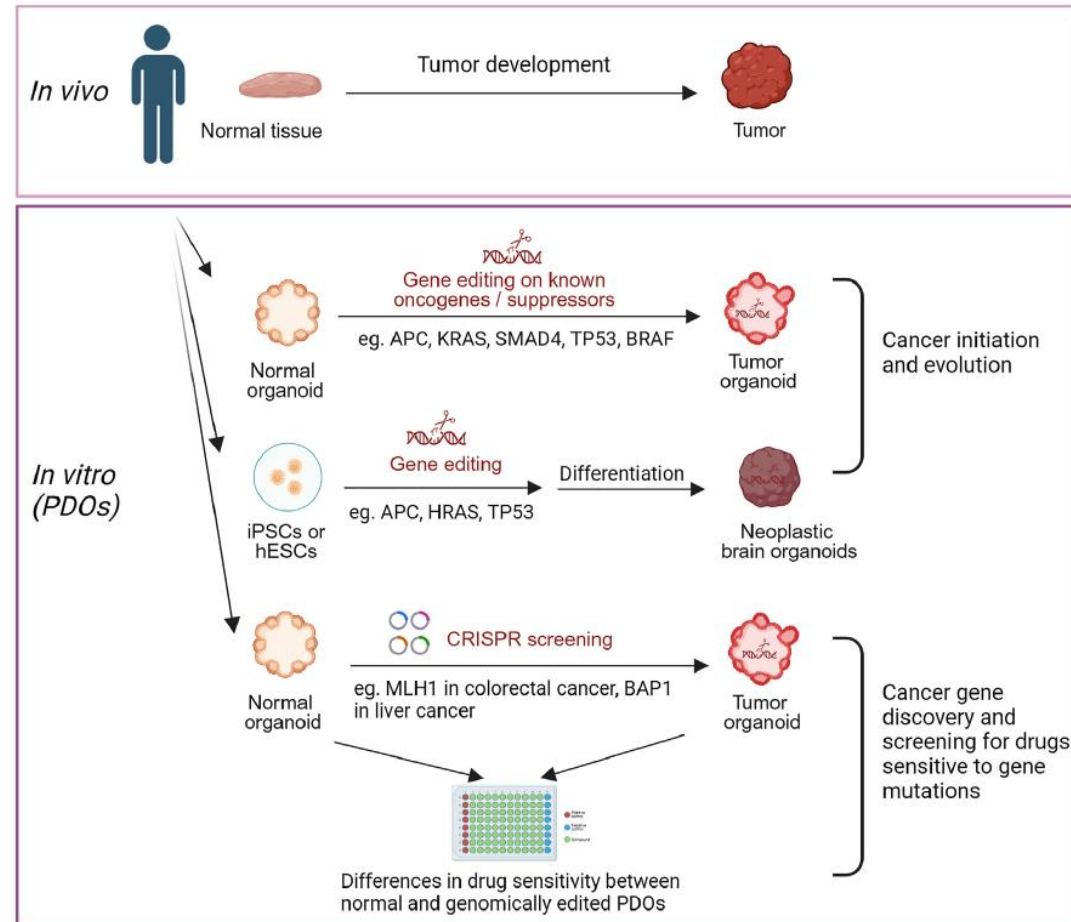
Patient-derived organoids (PDOs)



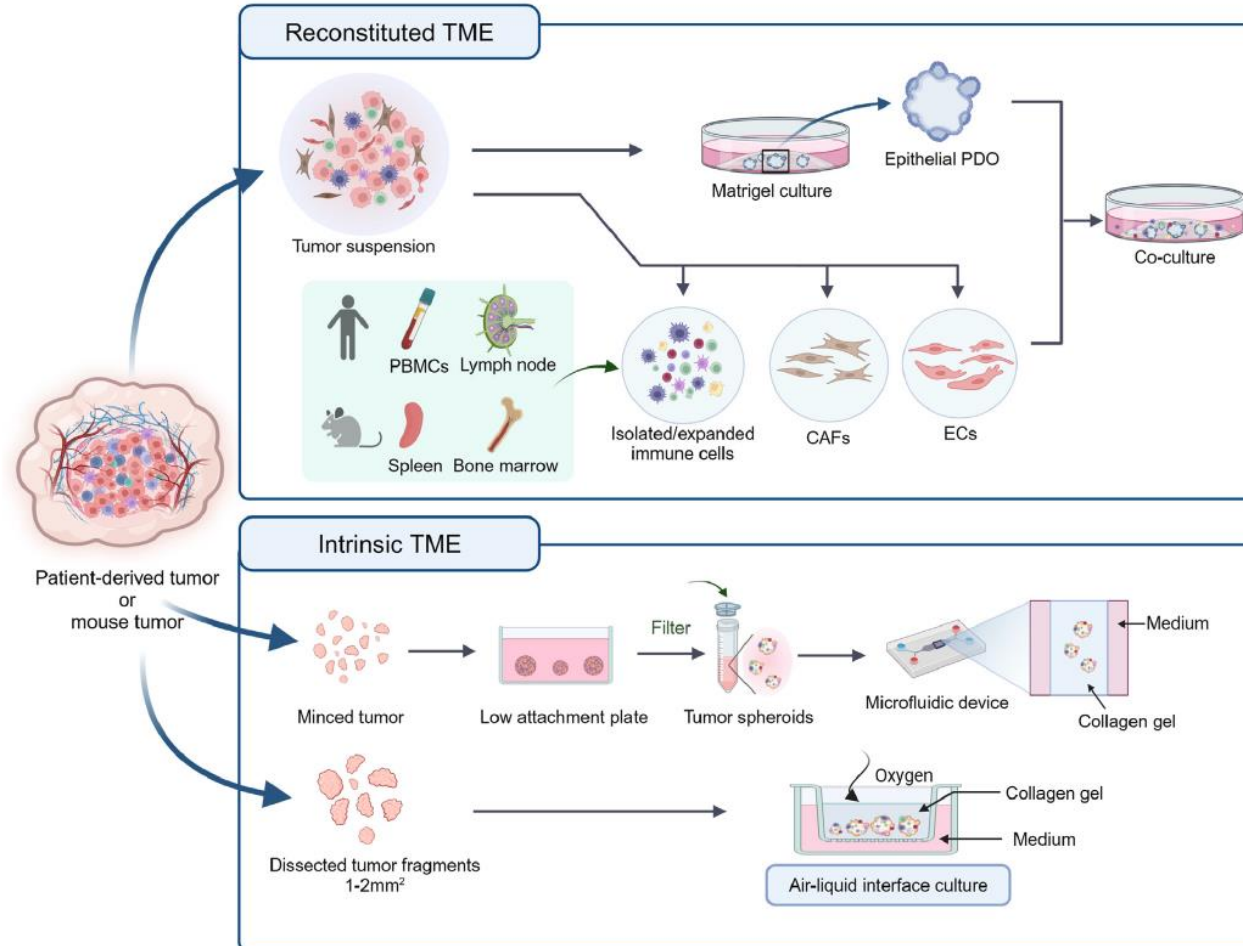
Patient-derived organoids (PDOs)



Patient-derived organoids (PDOs)



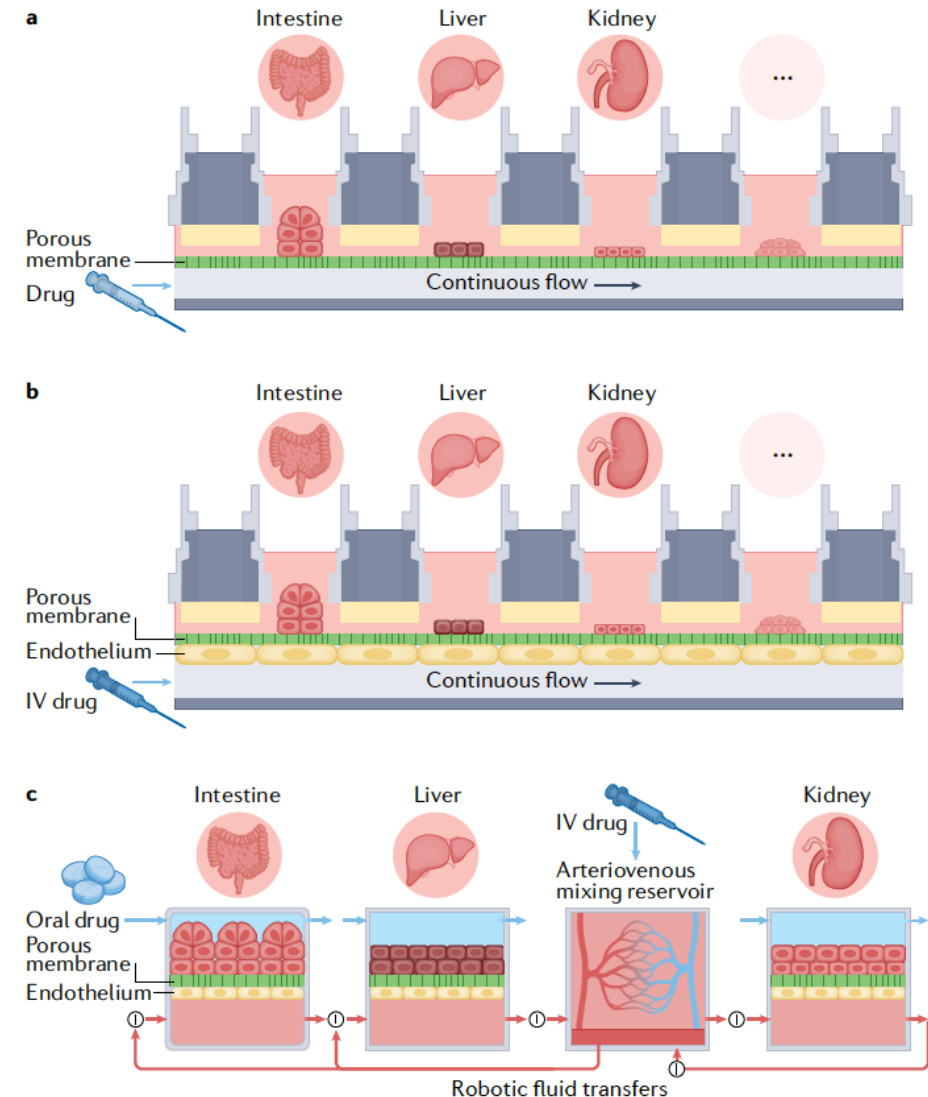
Patient-derived organoids (PDOs)



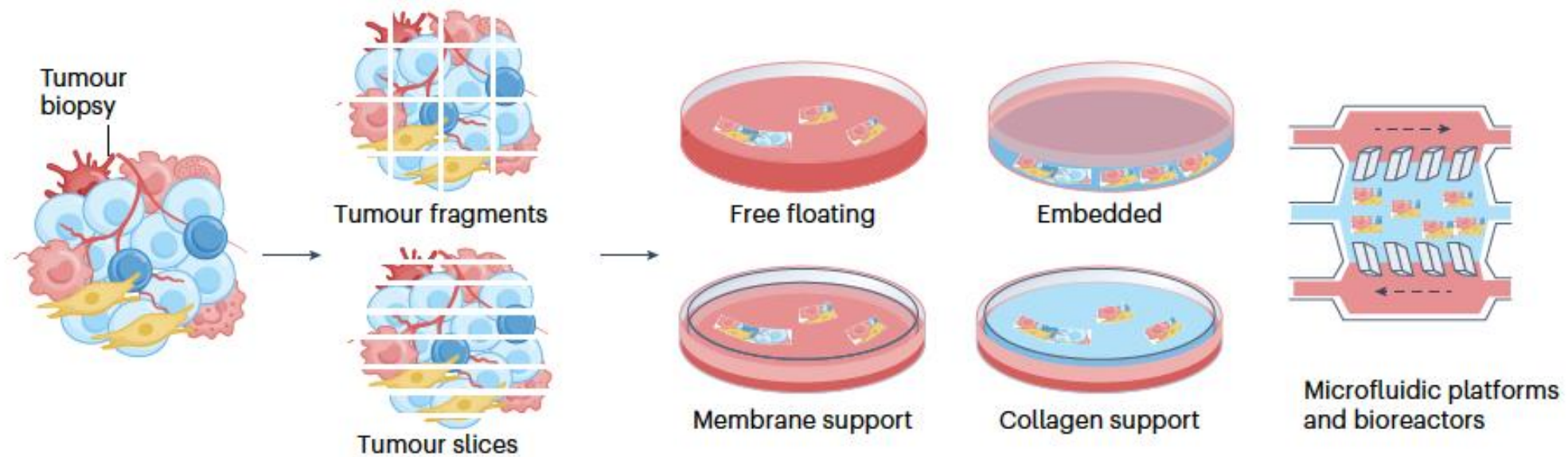
Organ-on-a-chip

Microfluidic culture devices that contain hollow channels lined by living cells and tissues cultured under dynamic flow

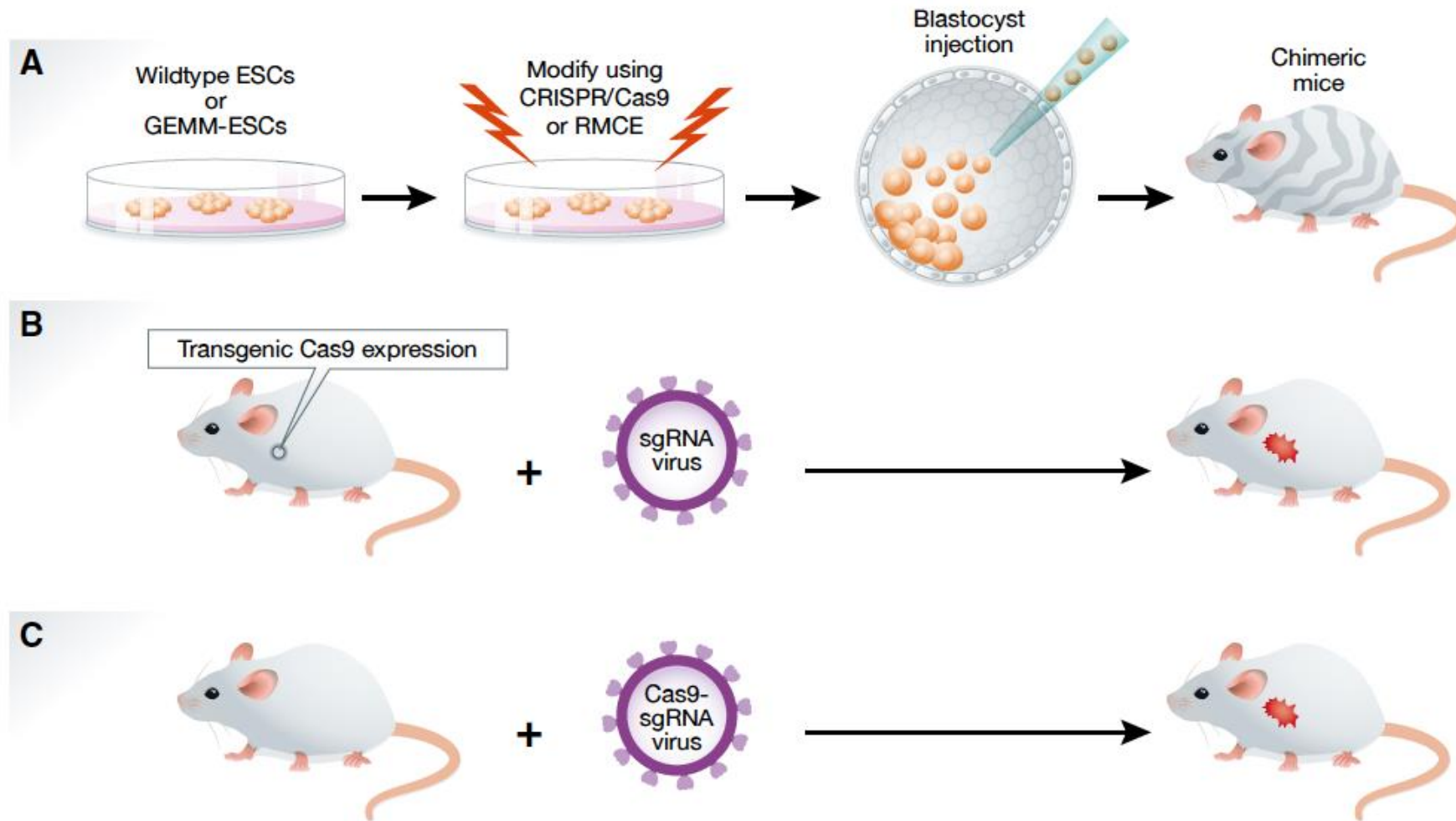
Fluid flow, shear stress, nutrient supply can be controlled



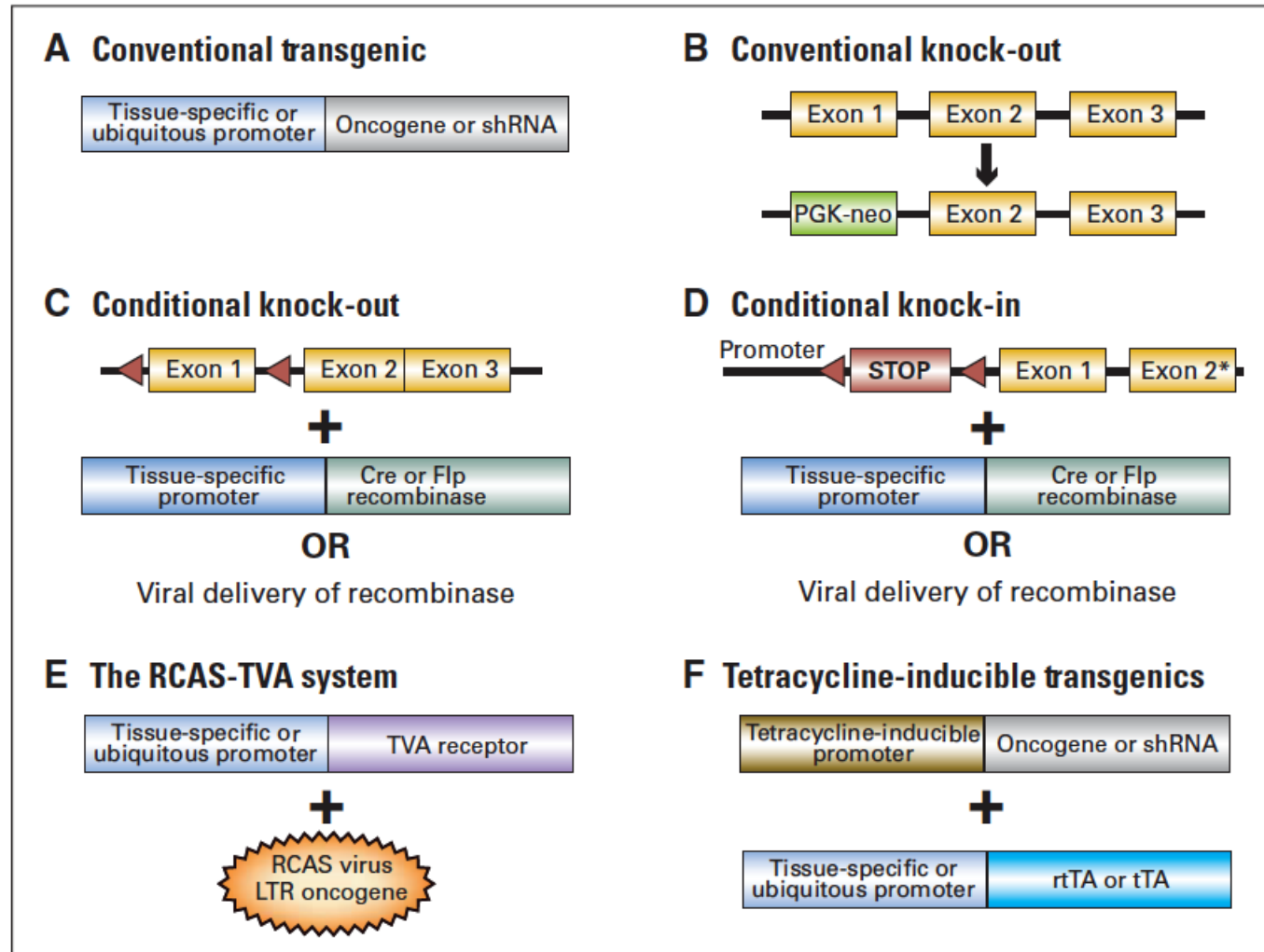
Non-organoid tumor explant culture



Genetically engineered mouse models (GEMMs)

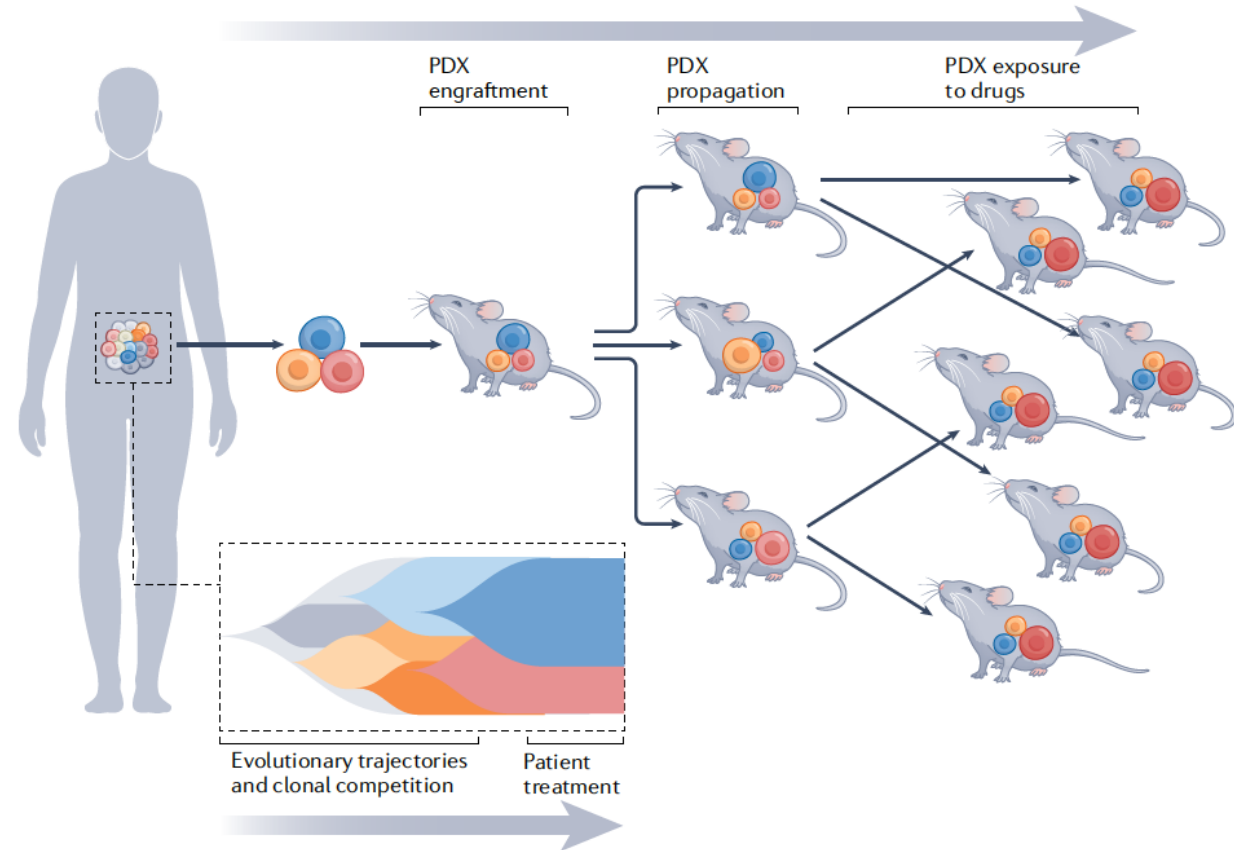


Genetically engineered mouse models (GEMMs)

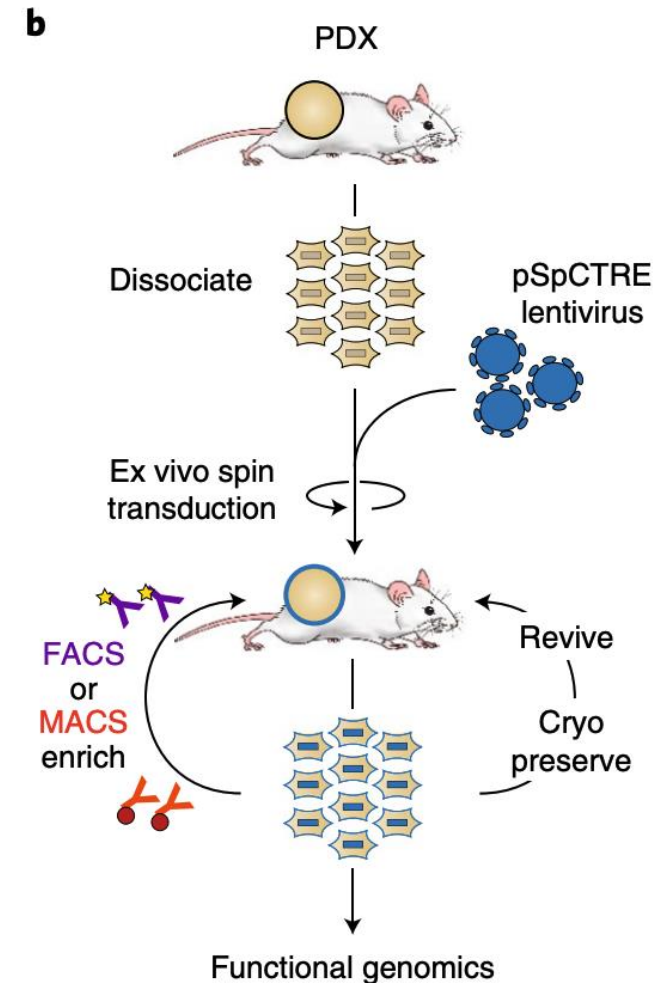
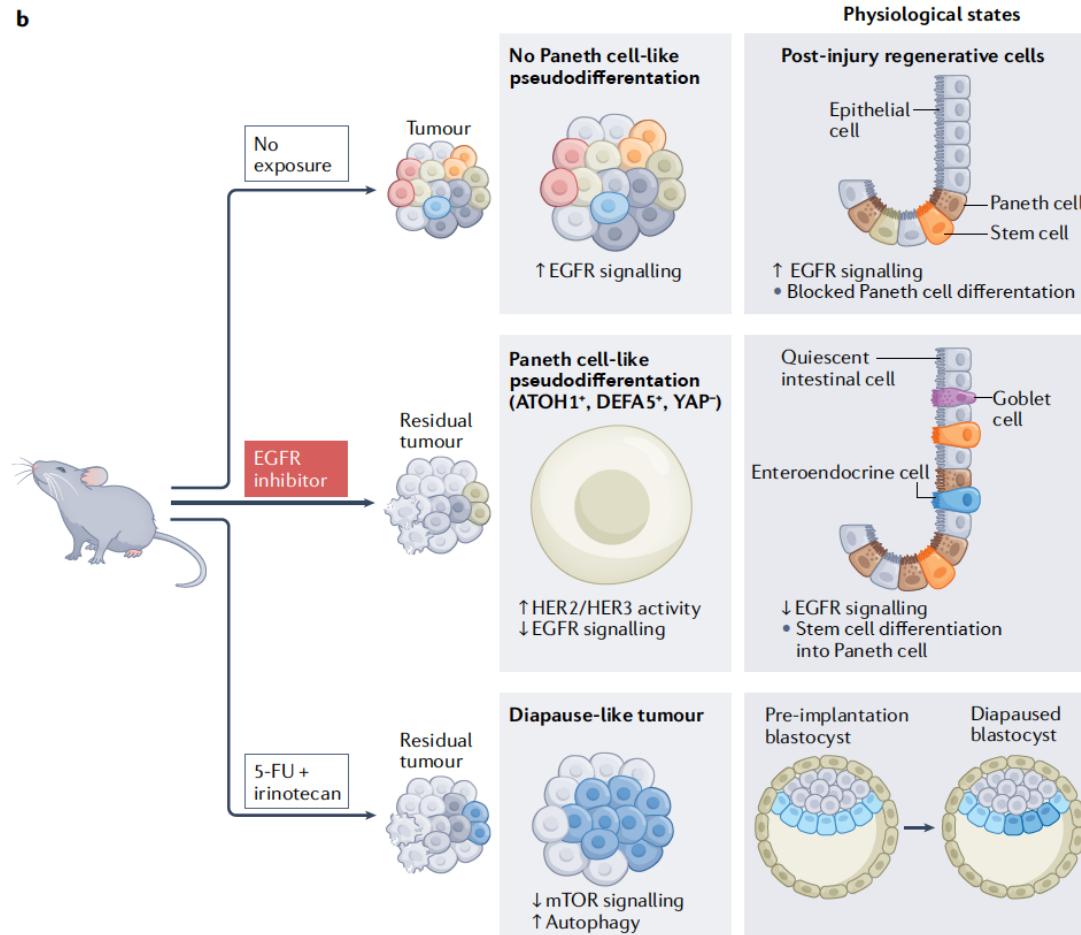


Patient-derived xenografts (PDXs)

Tumor fragments from surgical resections or biopsies that are propagated in mice through a series of implants and explants in immunodeficient mice



Patient-derived xenografts (PDXs)

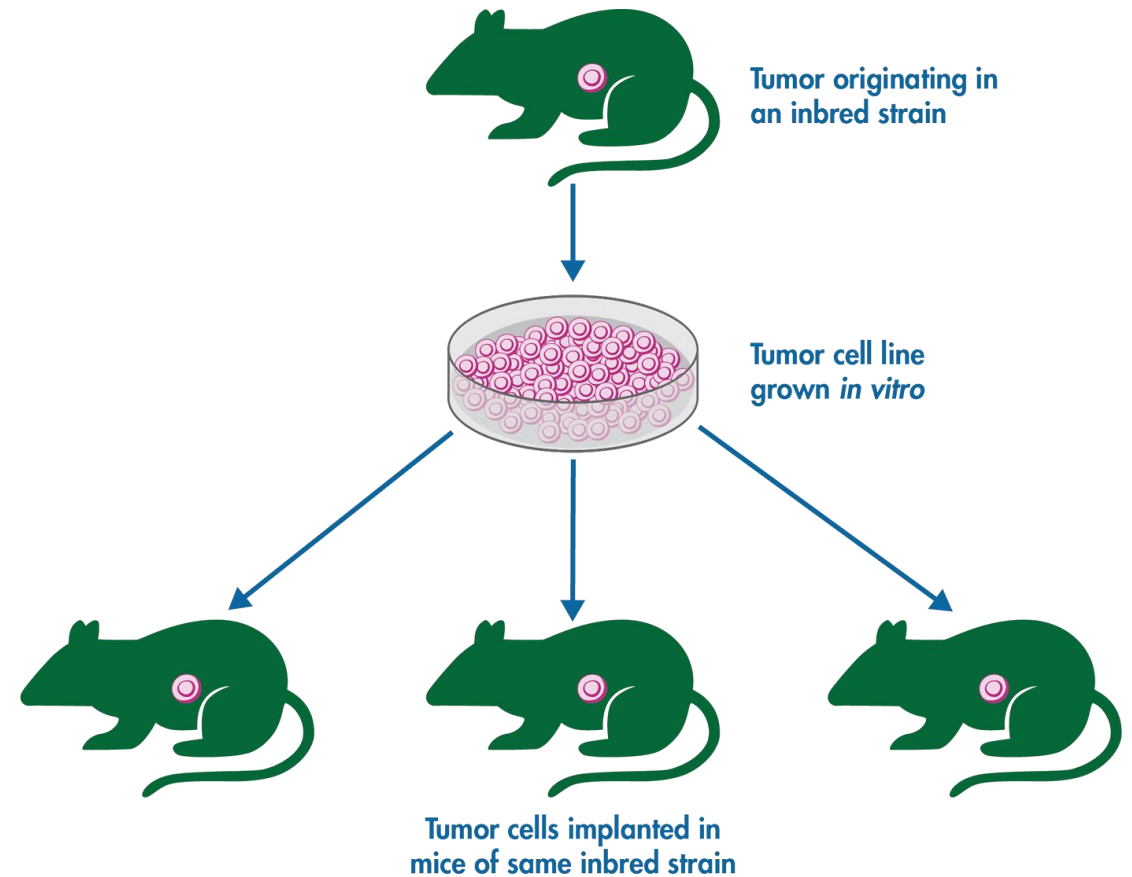


Syngeneic models

Models wherein cancer cells from one mouse are transplanted to another without the risk of rejection

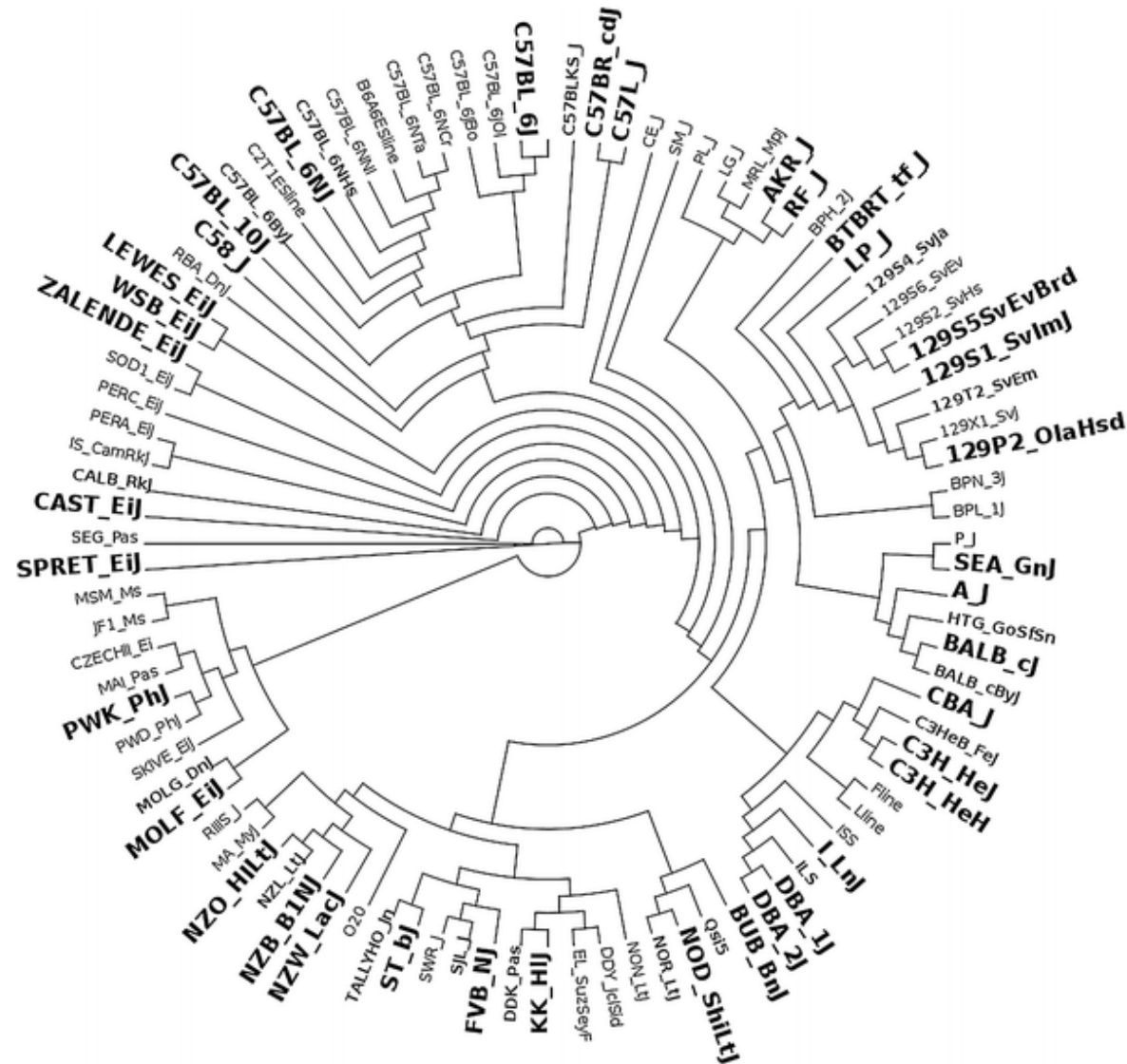
Immunocompetent – enabling immuno-oncology studies

Transplantation site is determined by the route of delivery (e.g. orthotopic vs subcutaneous)



How important is the site of transplantation?

A phylogenetic tree of inbred laboratory strains



Examples of immunodeficient transplant recipients

Strain Names	Mature B Cells	Mature T Cells	Dendritic Cells	Macrophages	Natural Killer Cells	Complement	Leakiness
NU/J (002019) Inbred Nude	Present	Absent	Present	Present	Present	Present	N/A
B6 Rag1 (002216) Rag1 KO <i>B6.129S7-Rag1^{tm1Mom}/J</i>	Absent	Absent	Present	Present	Present	Present	Absent
NOD scid gamma MHC Class I/II DKO (025216) NSG-MHC I/II DKO <i>NOD.Cg-Prkdc^{scid} H2-K1^{b-tm1Bpe} H2-Ab1^{g7-em1Mvw} H2-D1^{b-tm1Bpe} Il2rg^{tm1Wjl}/SzJ</i>	Absent	Absent	Defective	Defective	Absent	Absent	Absent

Quick Poll

You need to rapidly screen 50 different compounds for basic cell death. Which model would you start with?

- A) Cell line
- B) PDX mouse
- C) Tumor organoid

You are studying a very rare tumor. Which model would you start with?

- A) Cell line
- B) PDX mouse
- C) Tumor organoid

Case Study

You discover a rare, aggressive form of lung cancer defined by a complex fusion event involving a chromatin regulator. You want to study if this specific genetic change drives resistance to standard chemotherapy and if so, its underlying mechanism.

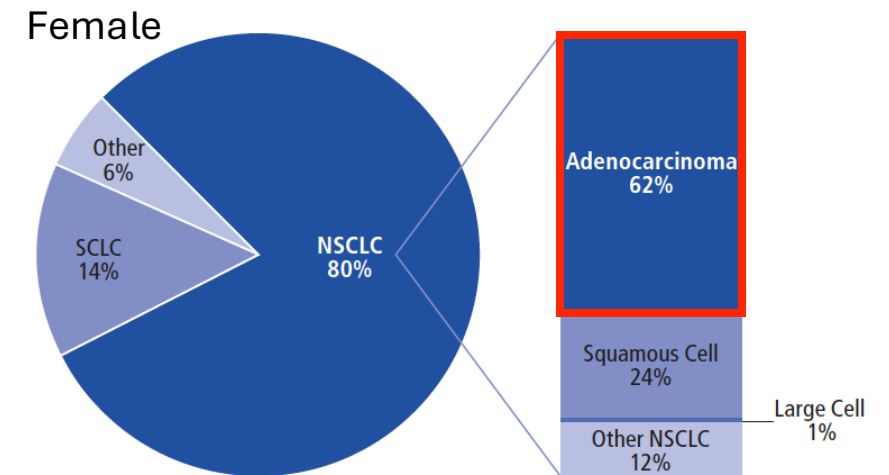
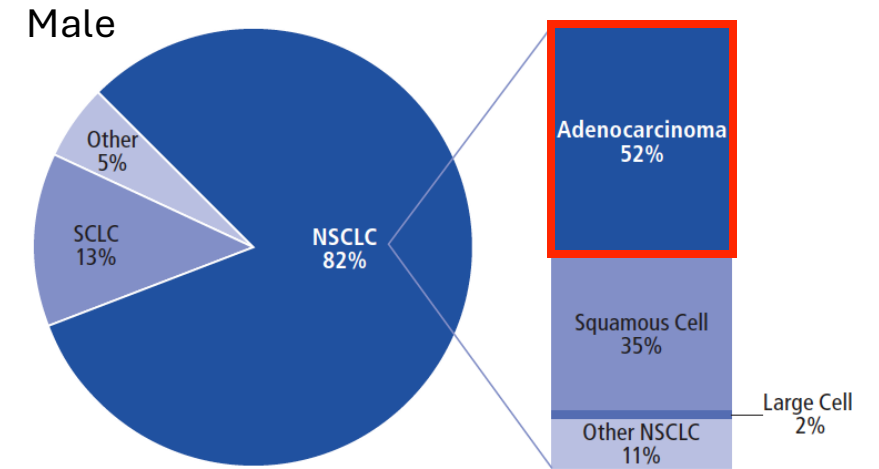
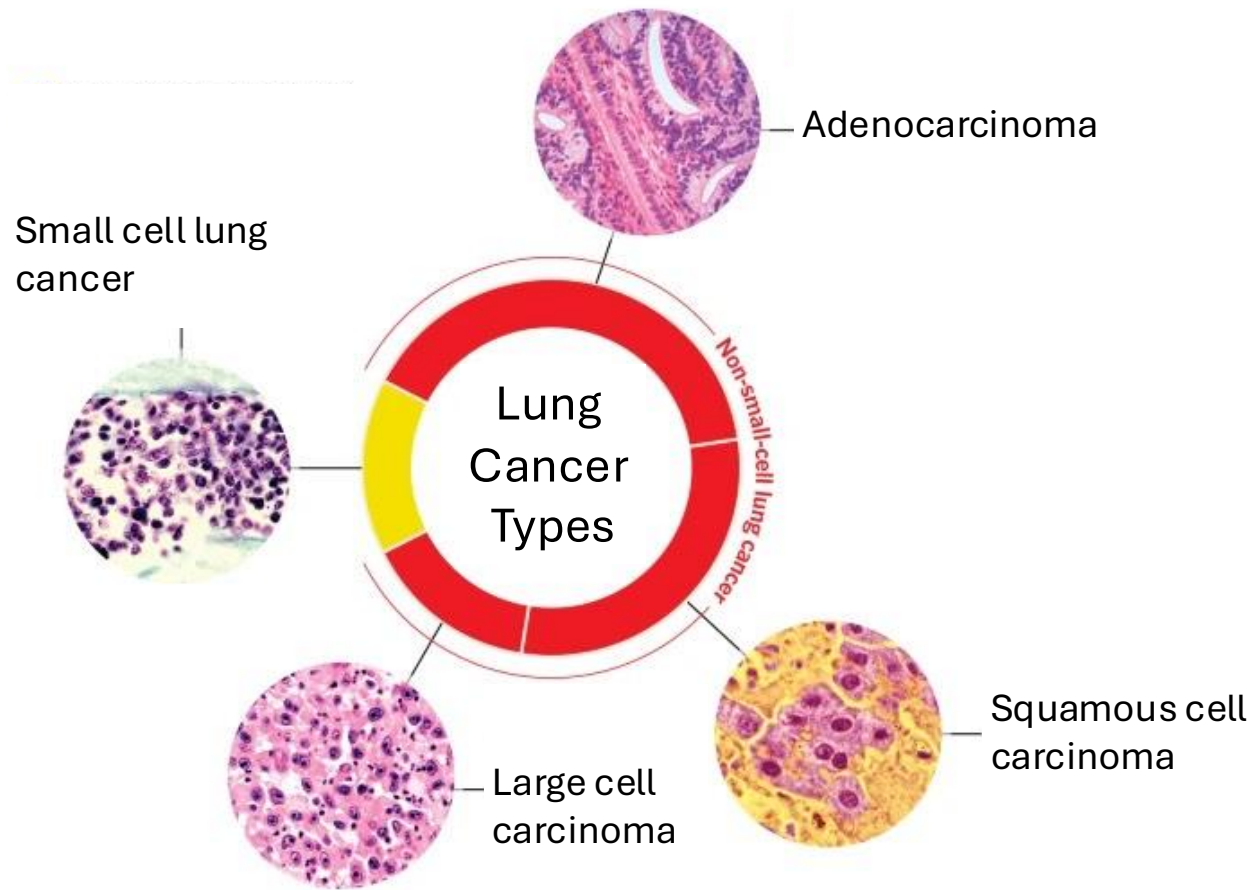
Cell Line Status: No commercial cell lines carry this specific fusion.

Patient Tissue: You have access to a fresh biopsy from one patient with this tumor.

Which preclinical model will you choose to determine if (and how) this genetic change drives resistance?

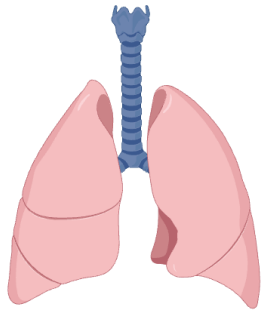
A (deeper) dive into the evolution of various lung cancer models

Types and subtypes of lung cancer

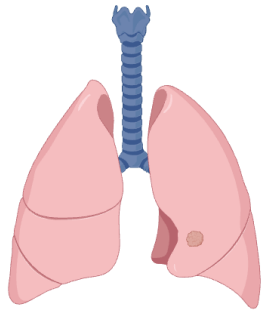


Lung cancer progression

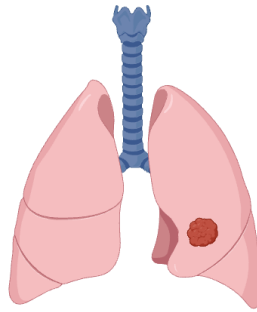
Normal



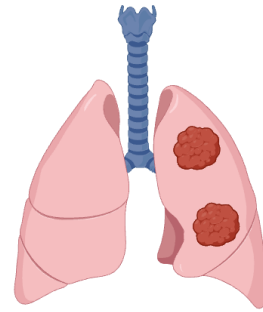
Pre-cancer



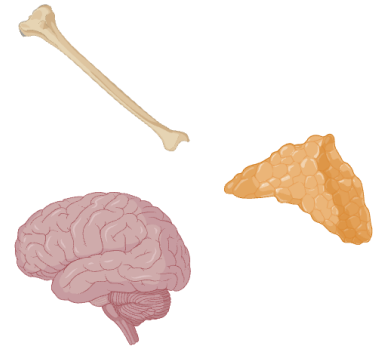
Early



Advanced



Metastasis

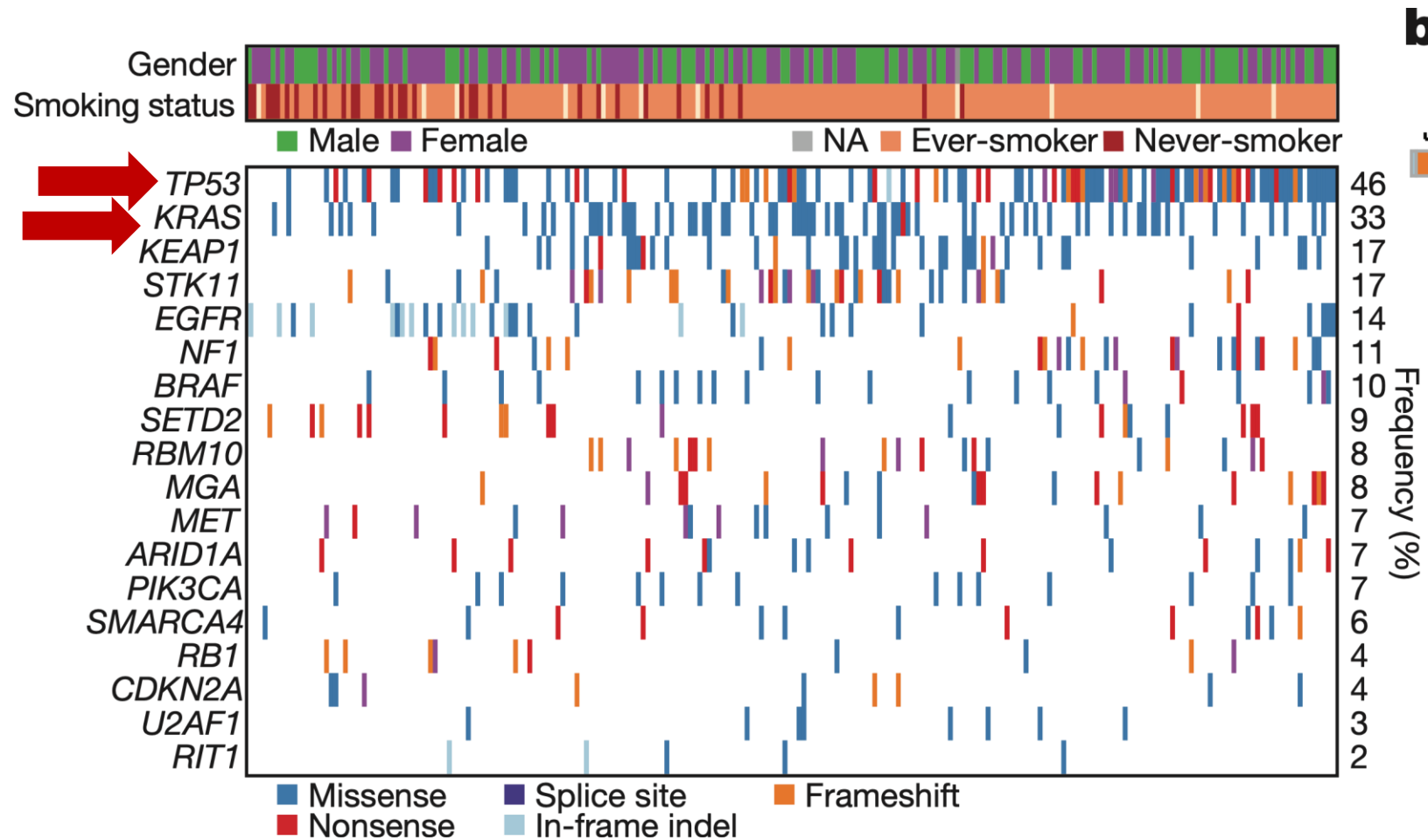


Smoking
Pollution
Occupational hazards
Other environmental factors

Tumor progression

Lineage fidelity

Somatic mutations in lung adenocarcinoma (LUAD)



The pioneer: the latent KRAS model ($Kras^{LA1}$ & $Kras^{LA2}$)

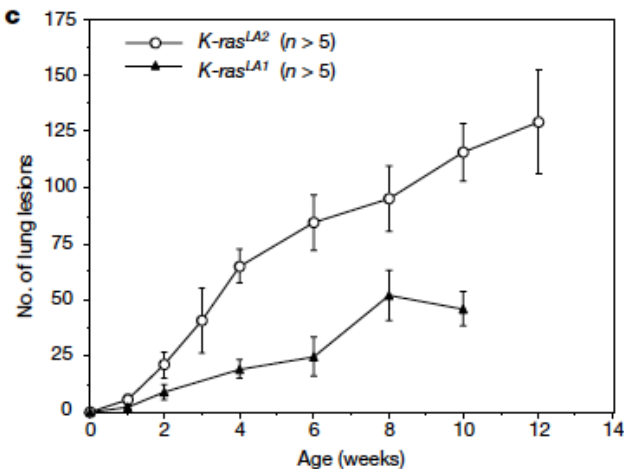
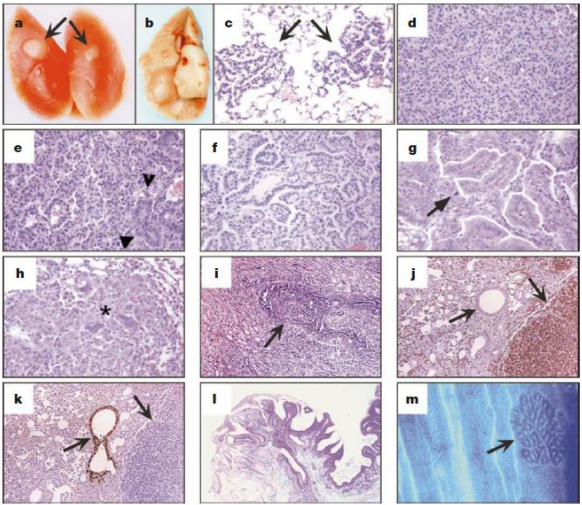
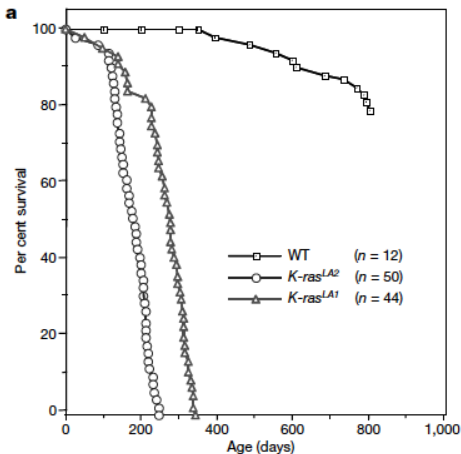
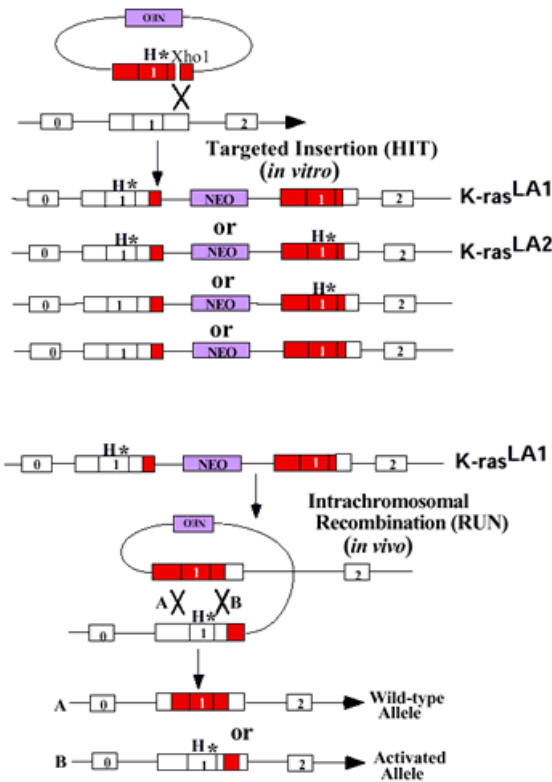
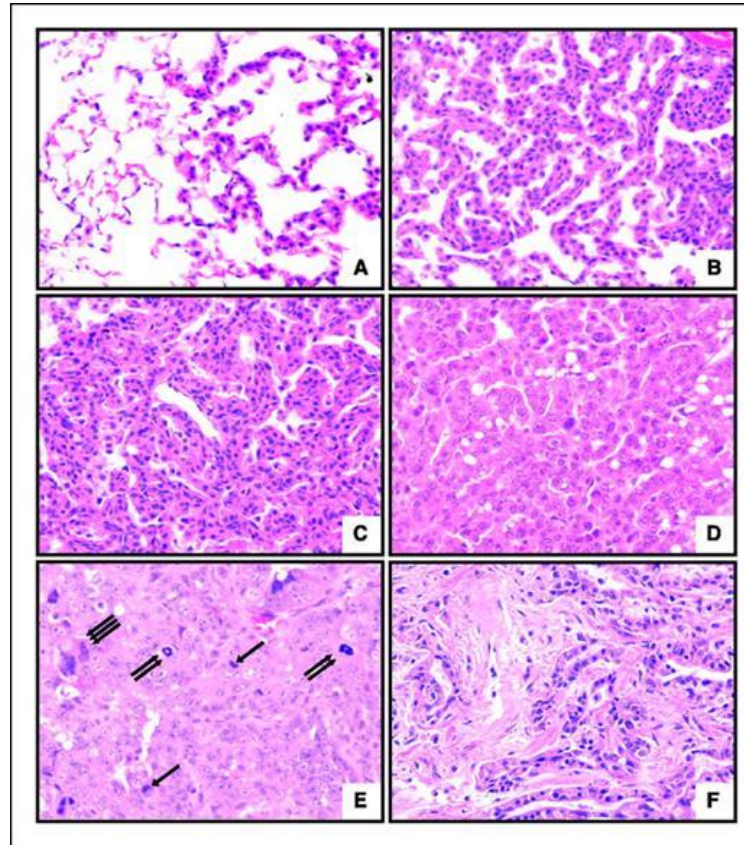
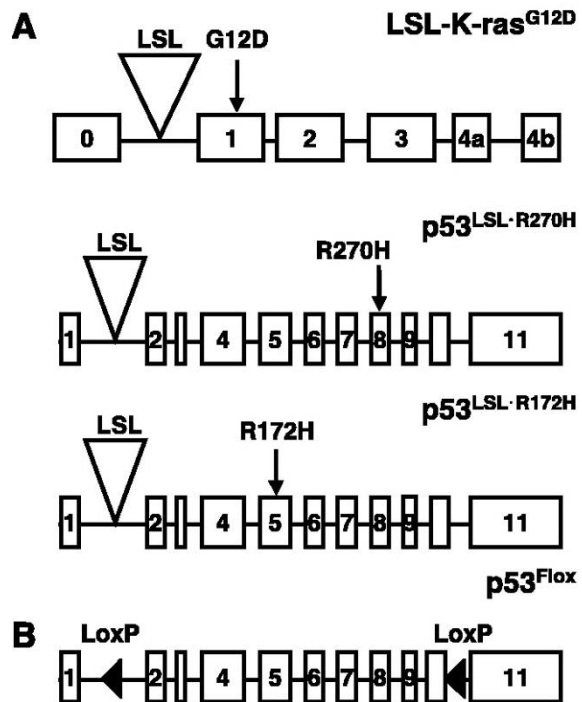


Table 1 Effect of *p53* status on the tumour spectrum in $Kras^{LA}$ mice

Tumour type	$Kras^{LA1/+}; p53^{+/-}$	$Kras^{LA1/+}; p53^{-/-}$
Lung adenocarcinoma	100	100
Thymic lymphoma	37	39
Papilloma	24	4
Fibrosarcoma	3	28
Haemangiosarcoma	2	33
Duodenal adenocarcinoma	0	2
Undifferentiated sarcoma	1	0
Histiocytic sarcoma	1	4
Medullablastoma	0	2
Osteosarcoma	0	2
Teratoma	0	2

Percentage of animals with a given tumour type. Numbers were combined for animals from both $Kras^{LA1}$ and $Kras^{LA2}$. $Kras^{LA1/+}; p53^{+/-}$ (n = 54); $Kras^{LA1/+}; p53^{-/-}$ (n = 87).

Temporal and spatial control: the conditional KP model



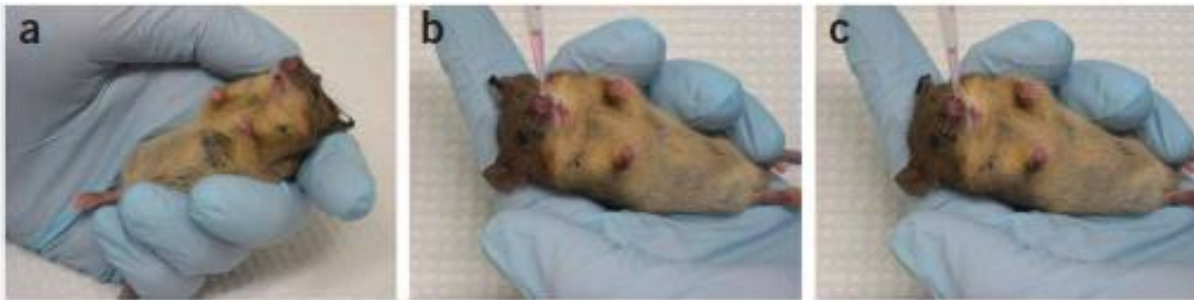
How to deliver Cre into lung cells?

- Intratracheal or intranasal
 - Adenovirus-Cre
 - Lentivirus-Cre
- Germline
 - Cell-type specific CreER mouse lines (e.g. *Sftpc-CreER*, *Scgb1a1-CreER*)

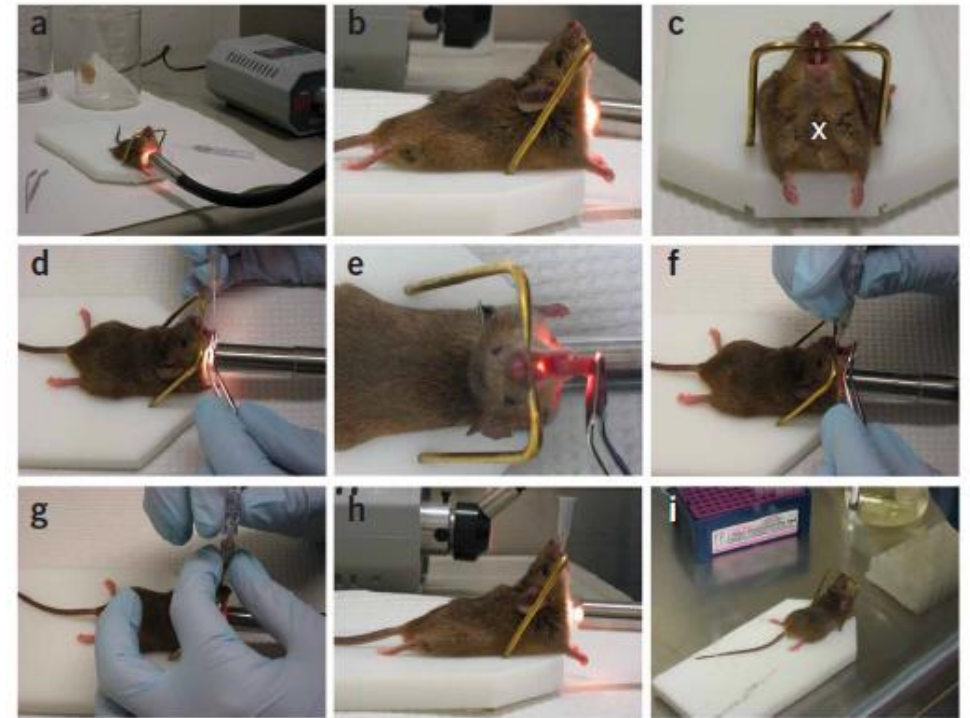
Temporal and spatial control: the conditional KP model

How to deliver Cre into lung cells?

Intranasal



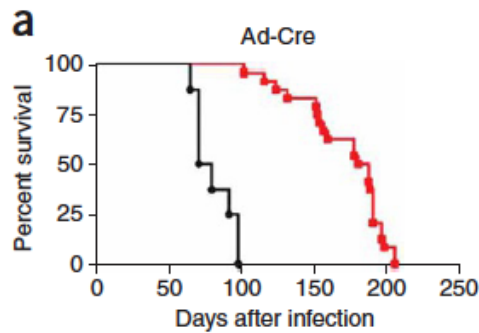
Intratracheal



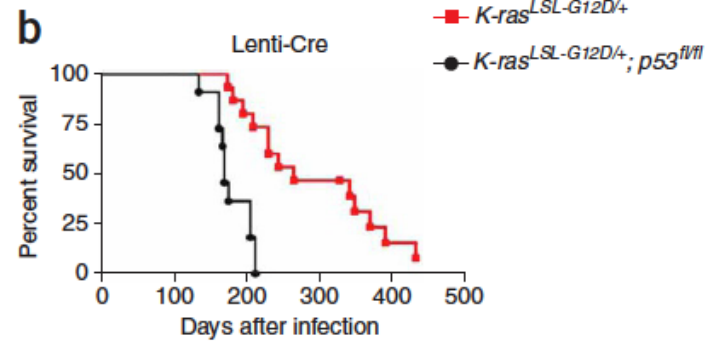
Temporal and spatial control: the conditional KP model

How to deliver Cre into lung cells?

Adenovirus

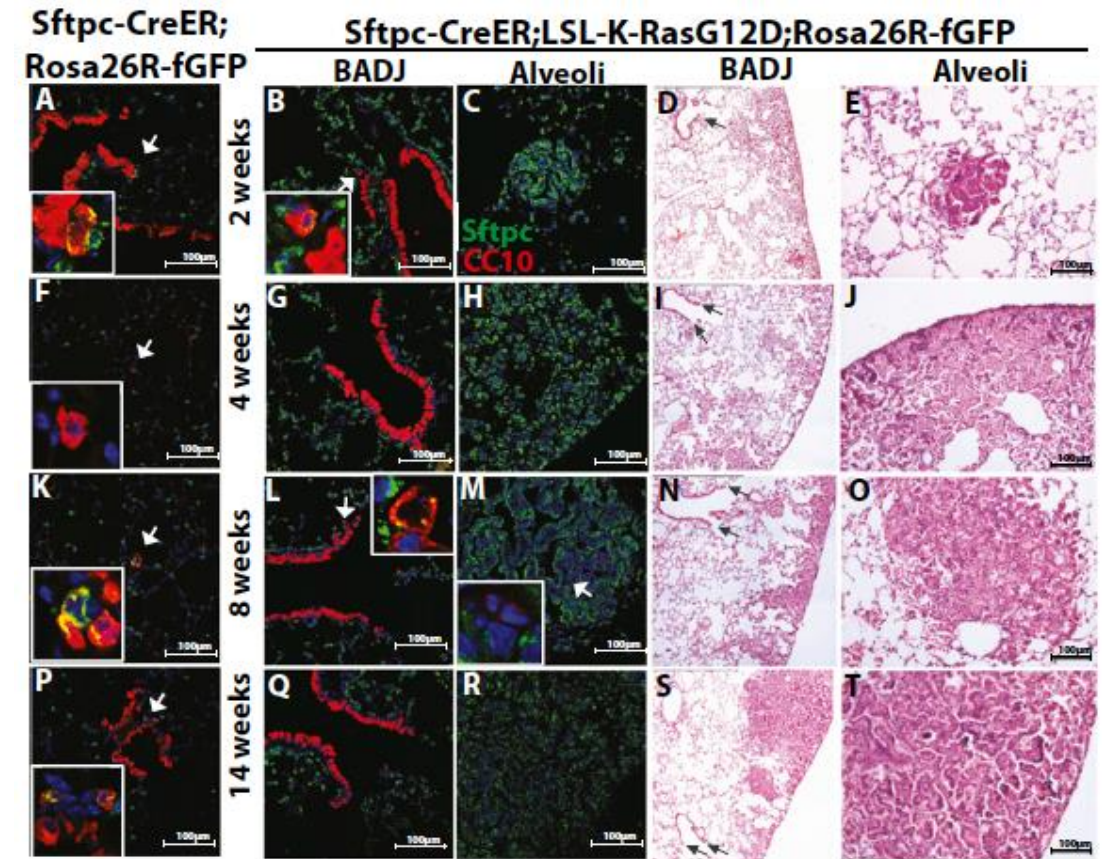


Lentivirus

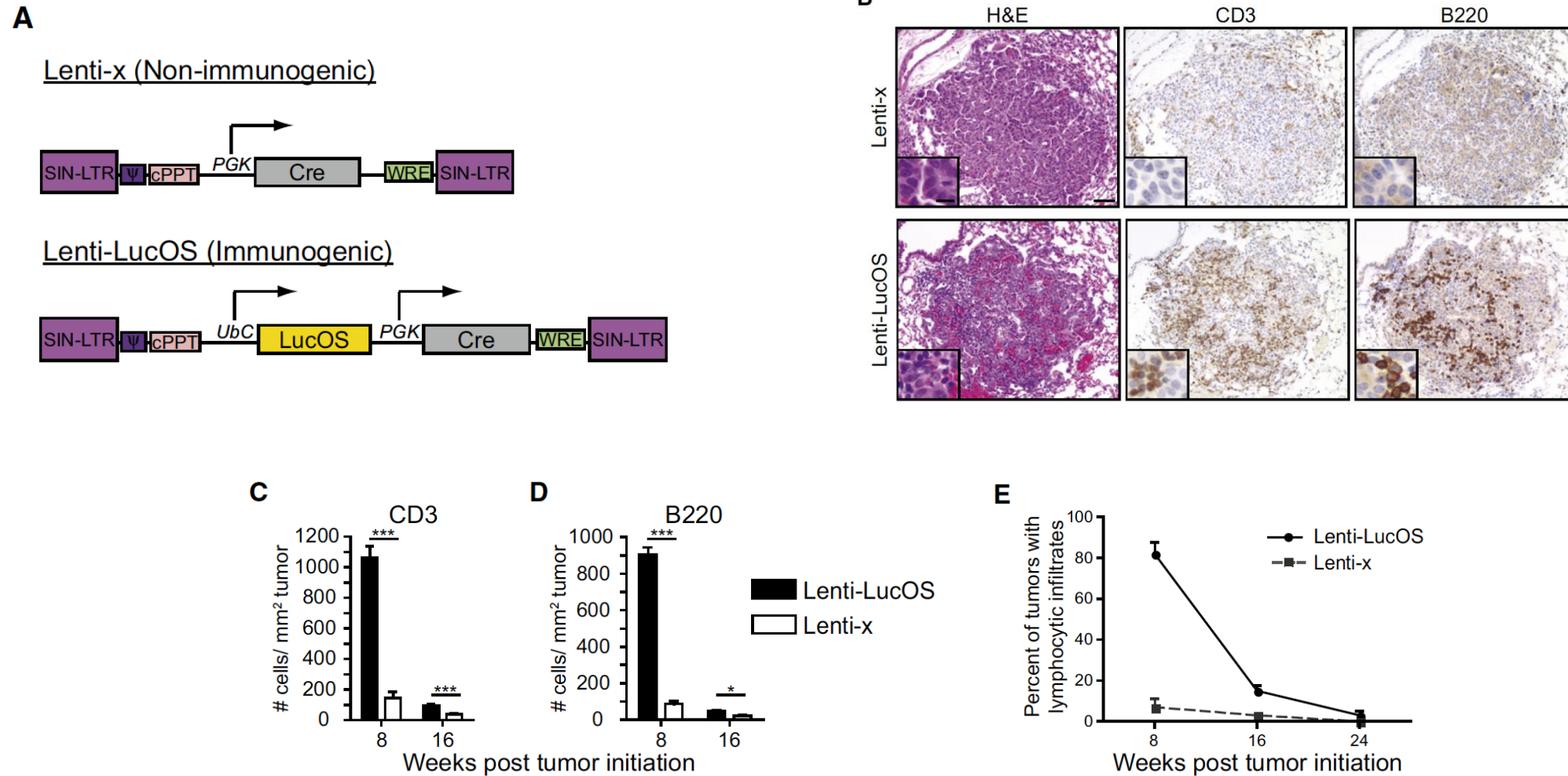


— $K\text{-ras}^{\text{LSL-G12D/+}}$
● $K\text{-ras}^{\text{LSL-G12D/+}; p53^{\text{fl/fl}}}$

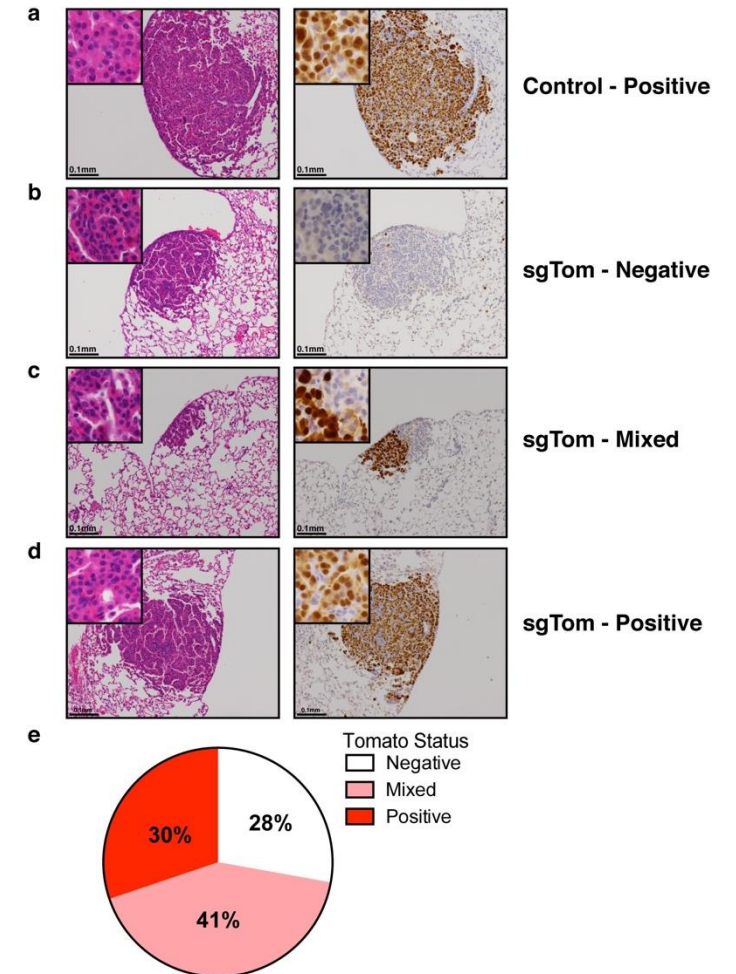
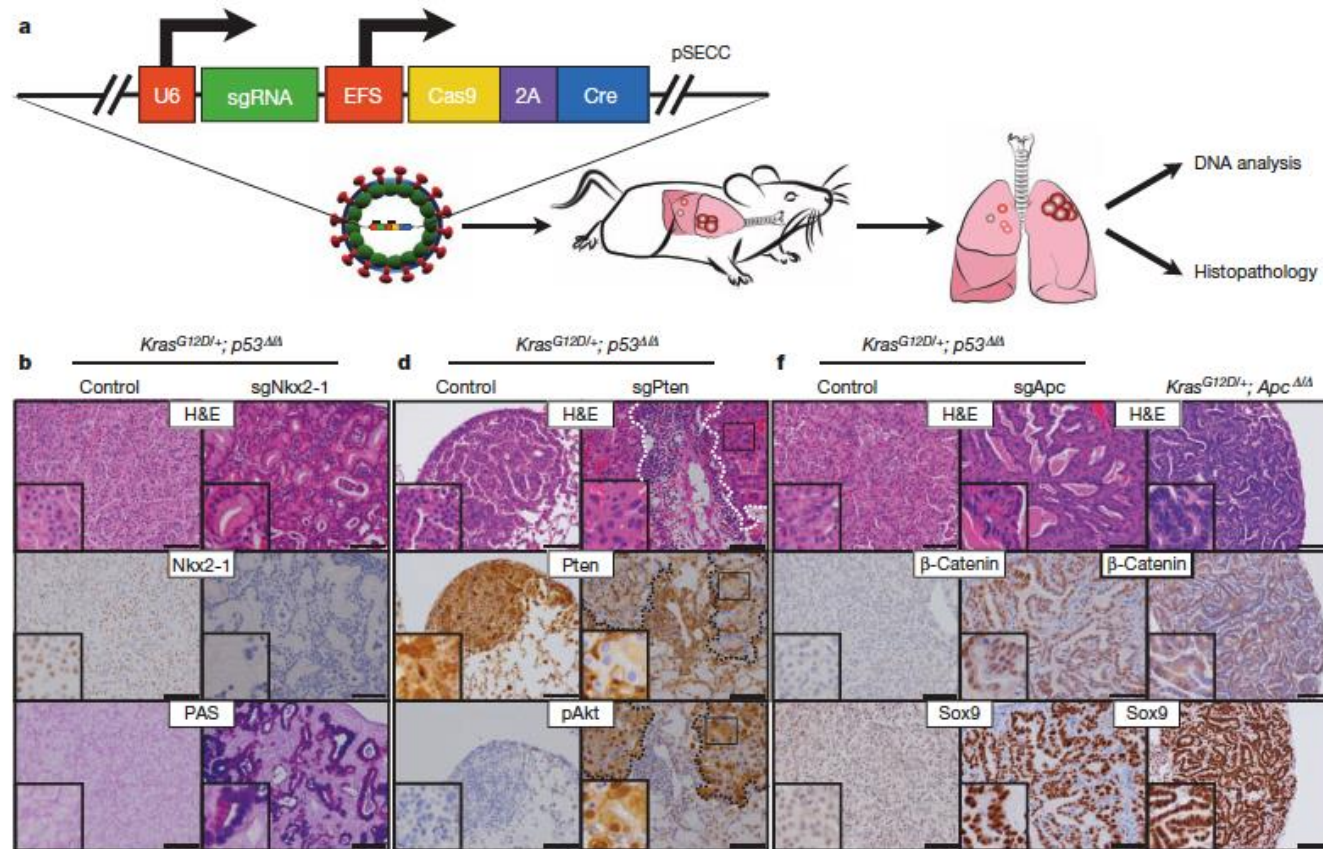
Germline



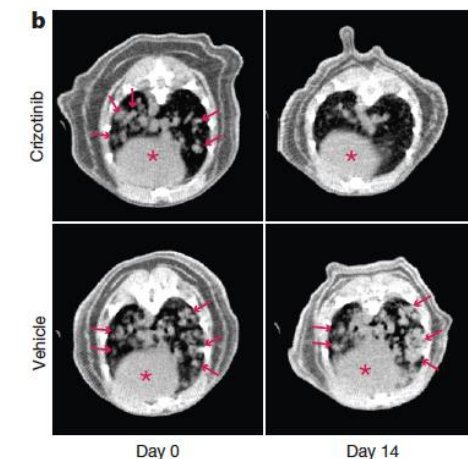
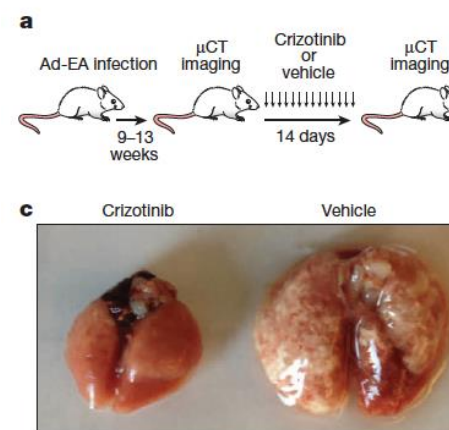
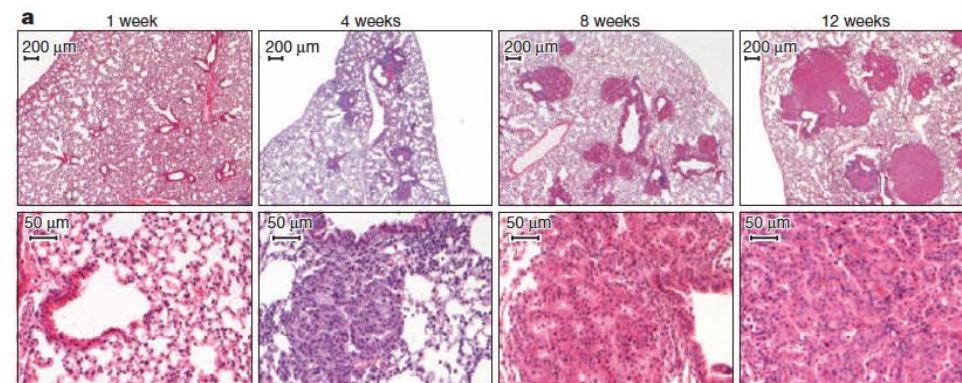
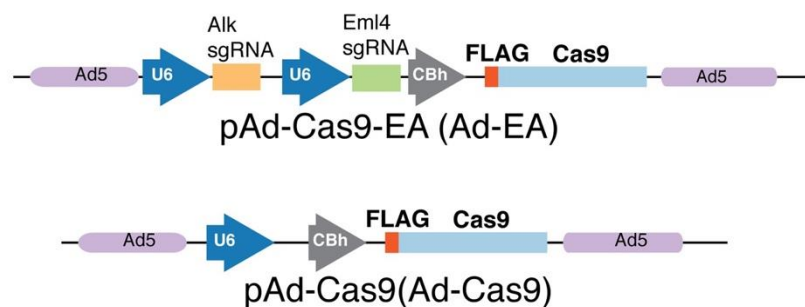
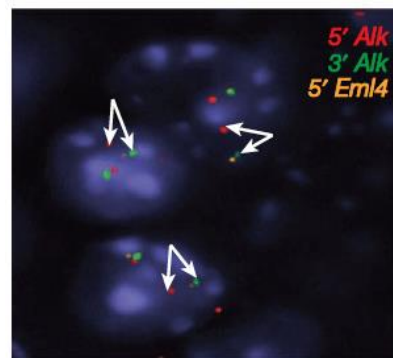
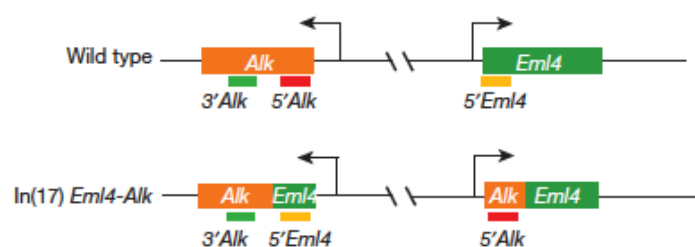
Immunogenic KP models



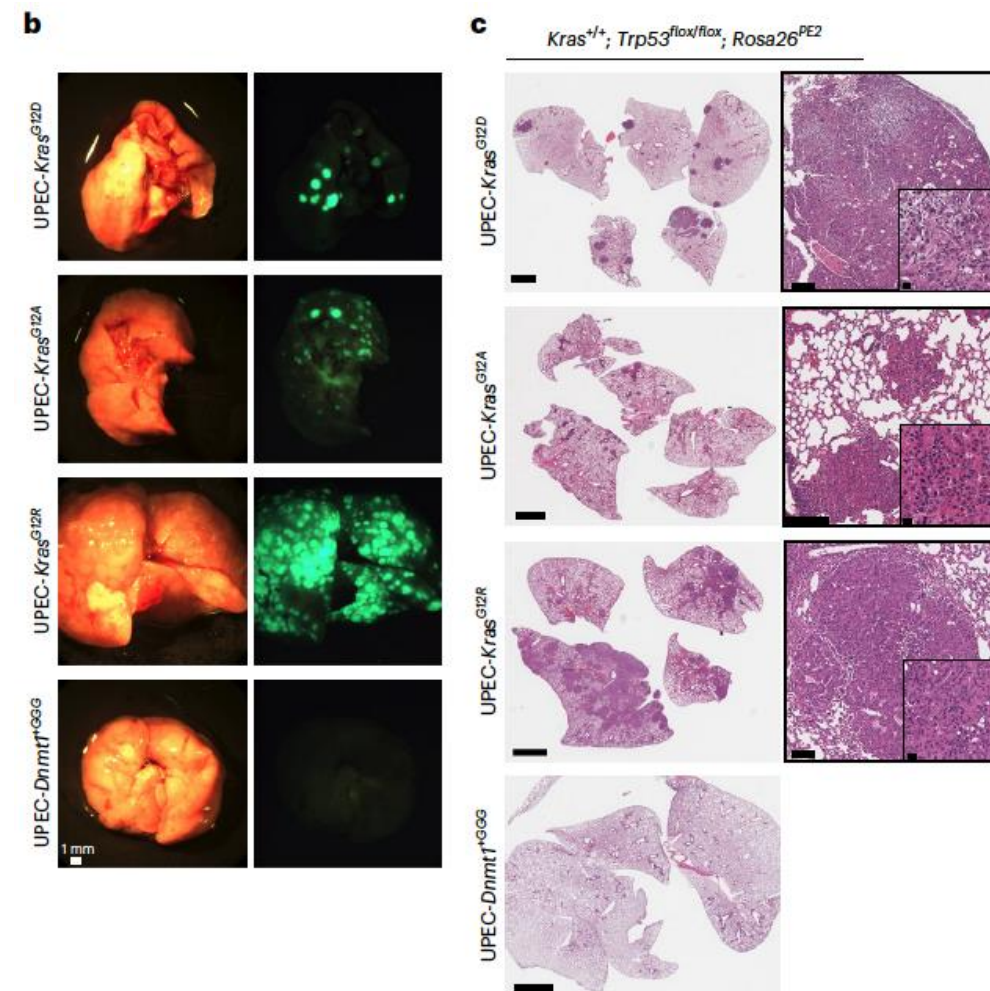
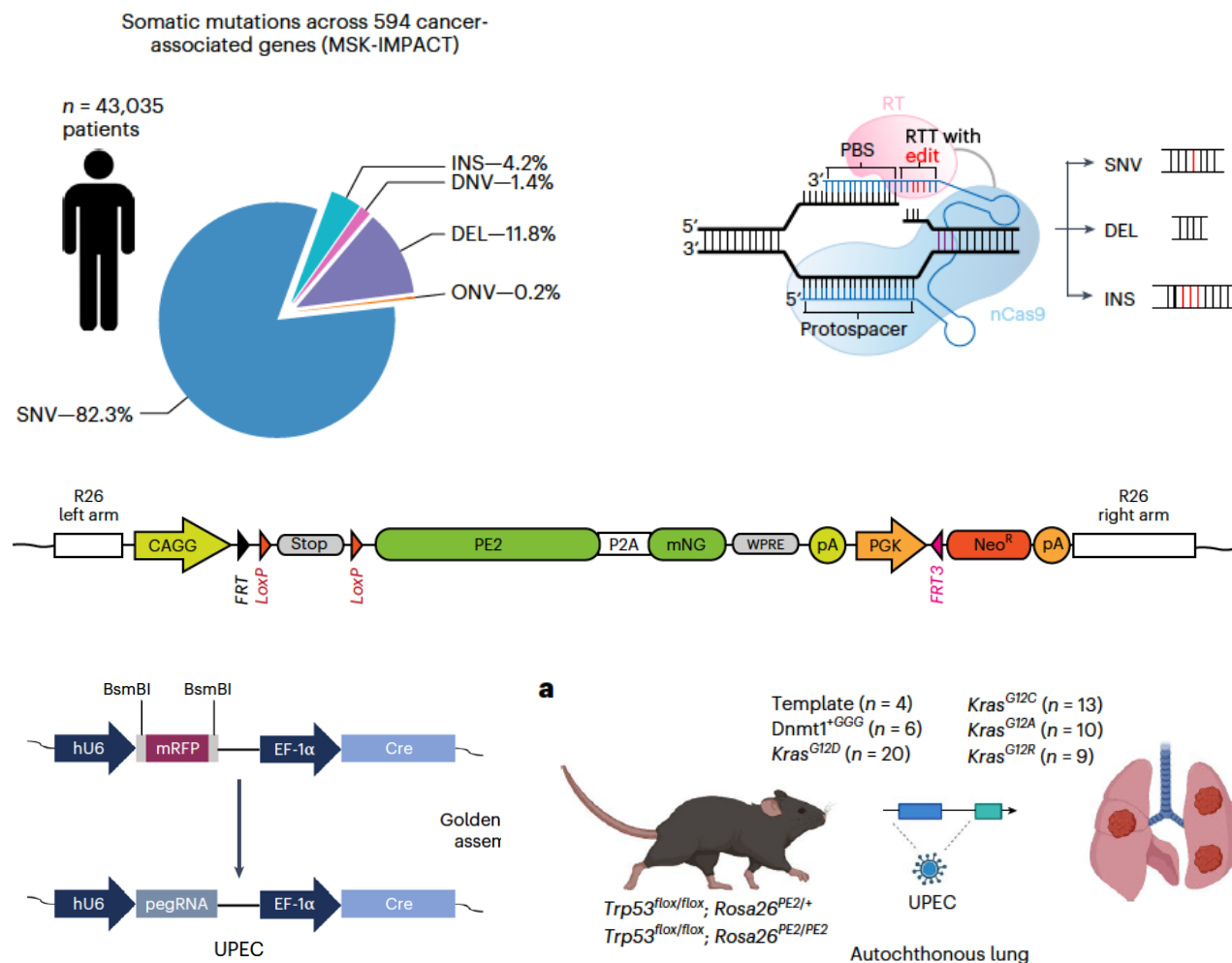
KP models in the CRISPR era



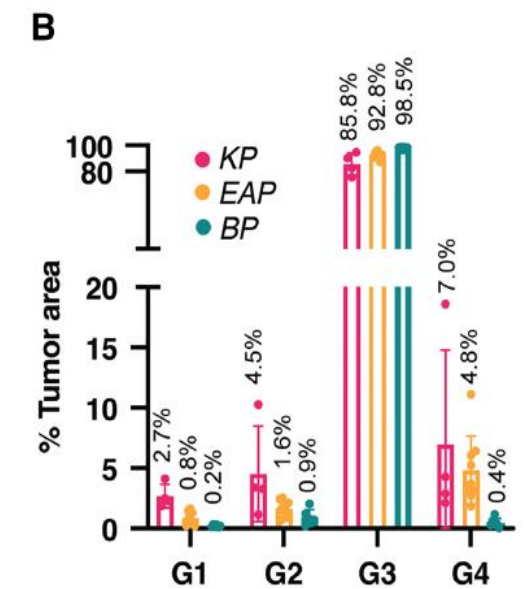
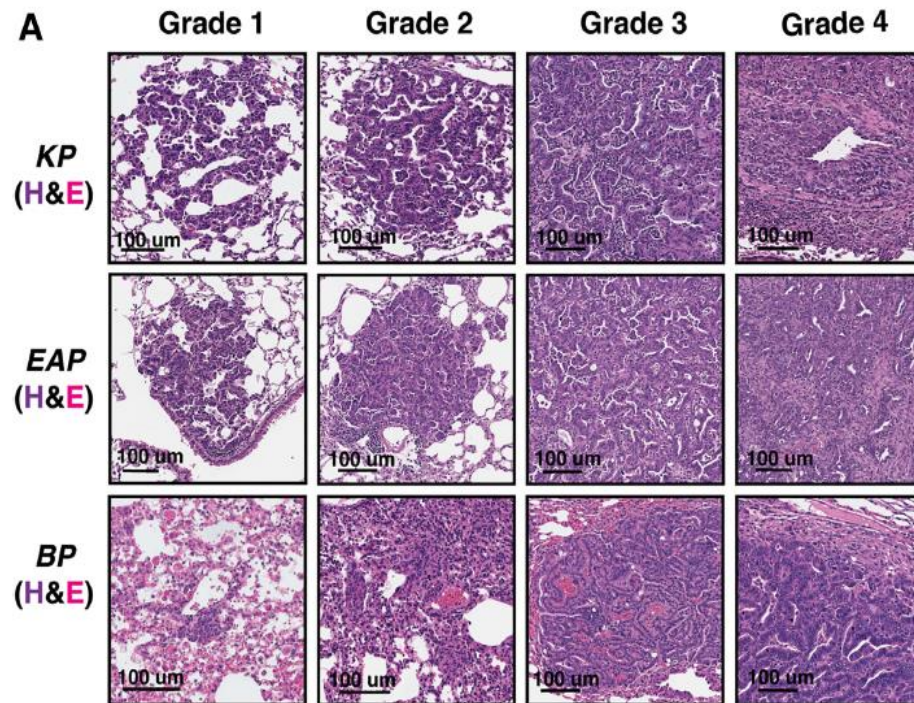
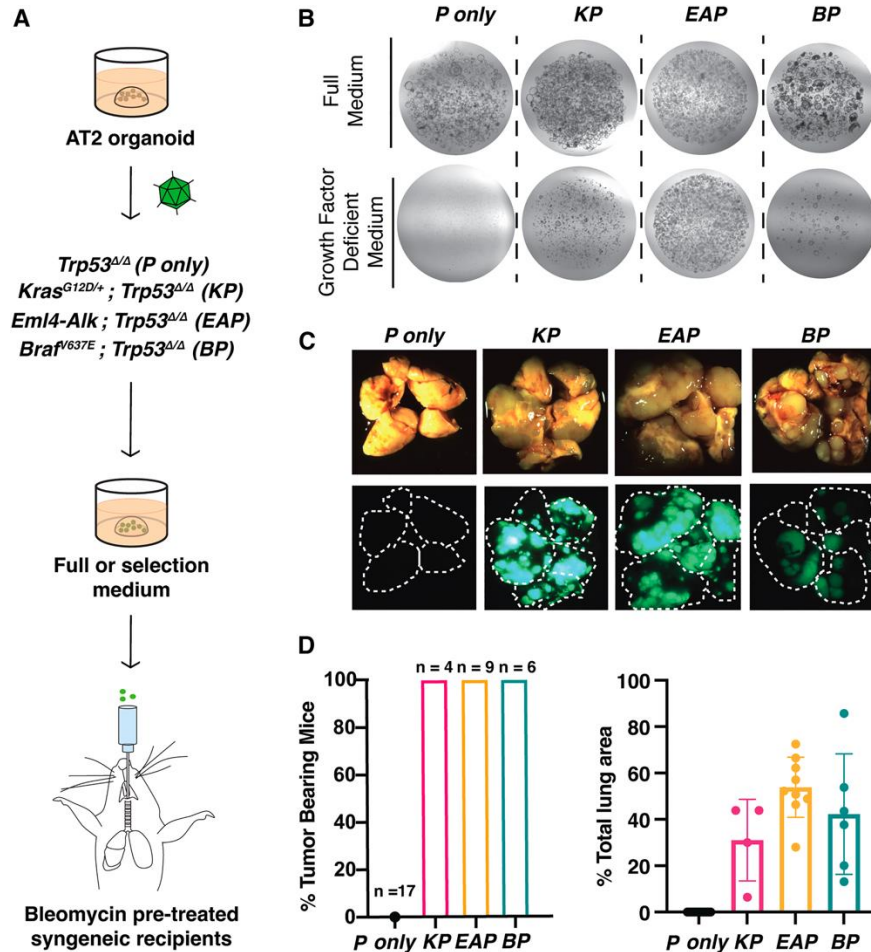
Lung models via CRISPR beyond KP



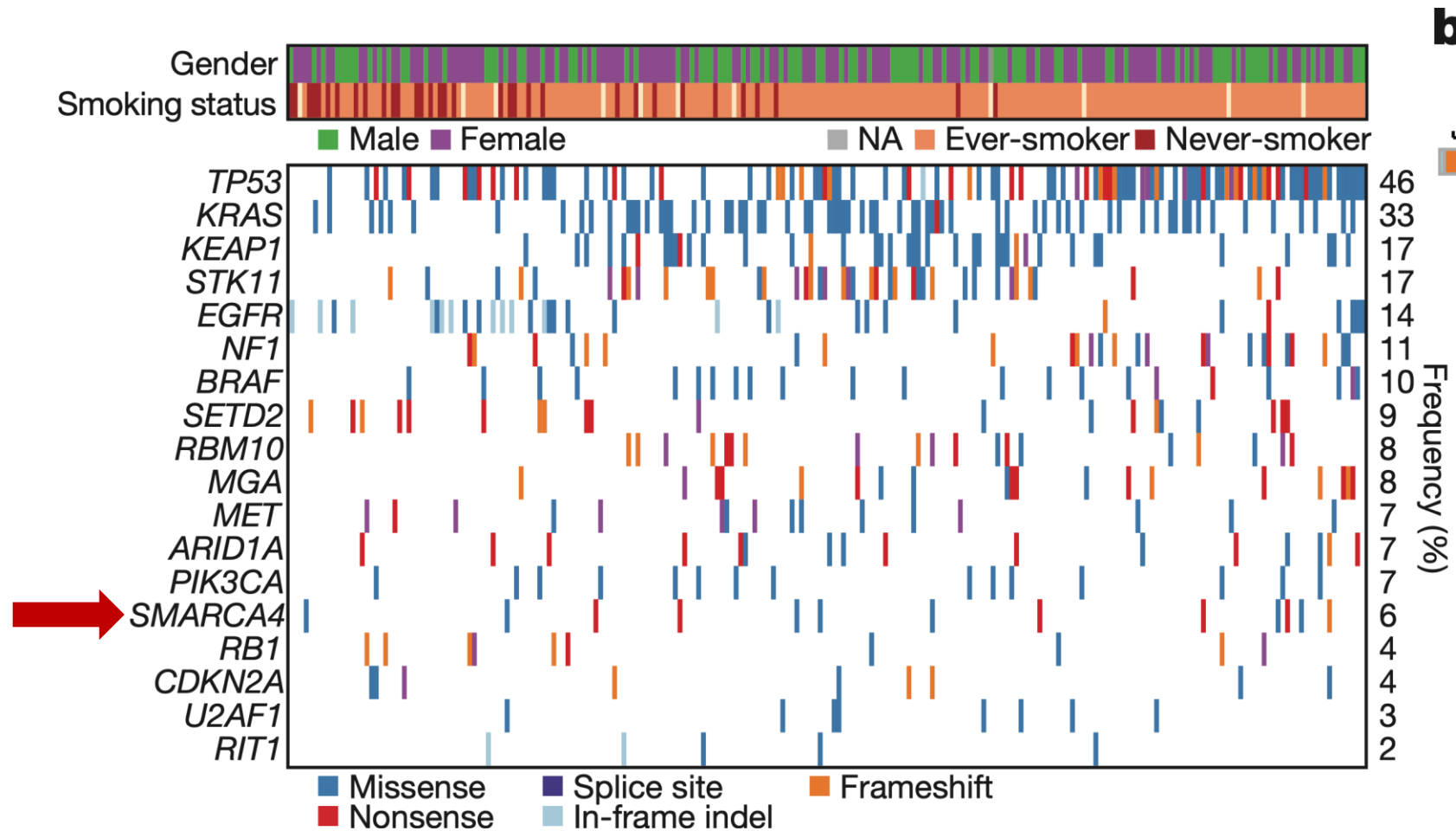
KP models in the precision genome editing era



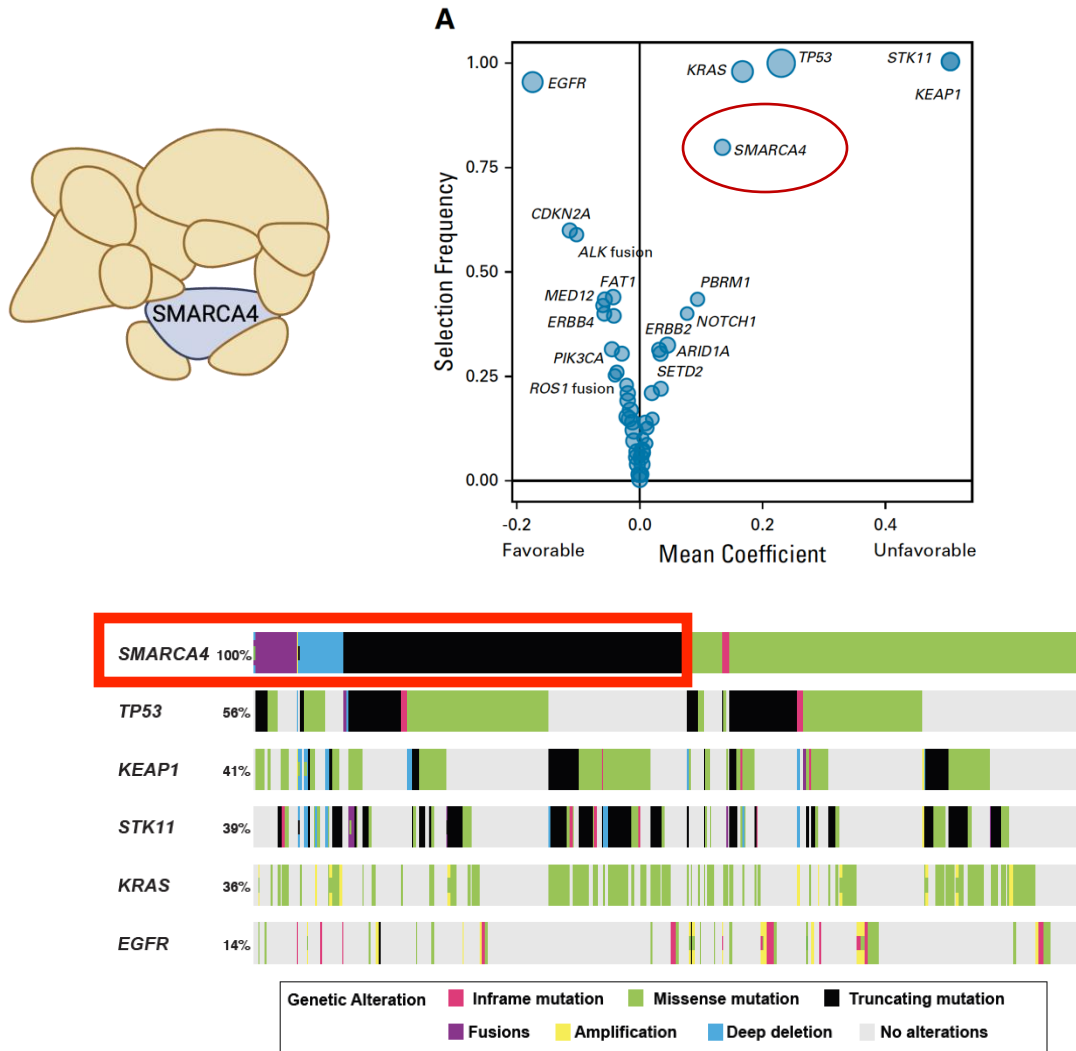
Don't forget about organoids!



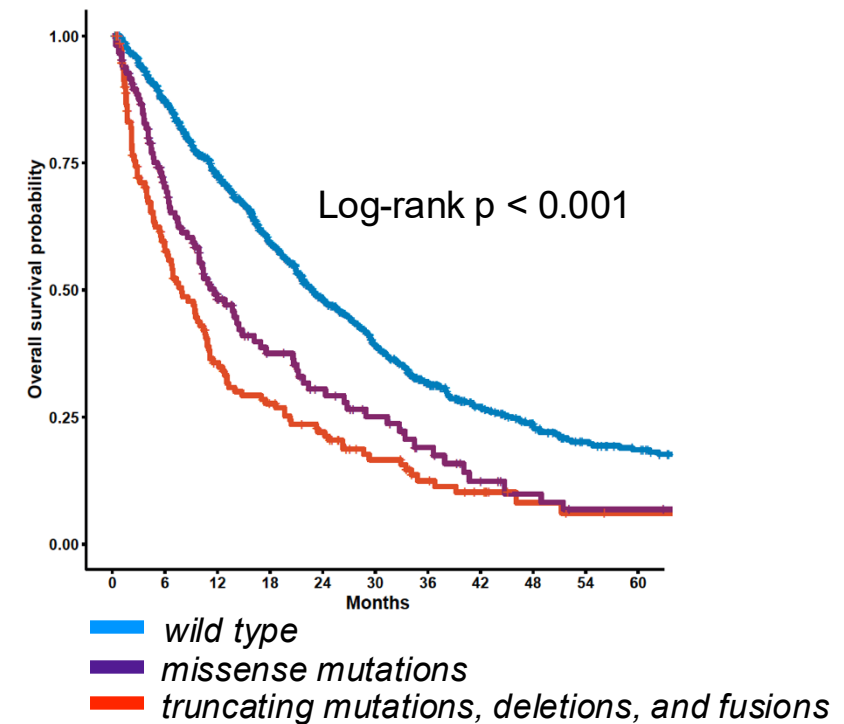
Somatic mutations in lung adenocarcinoma



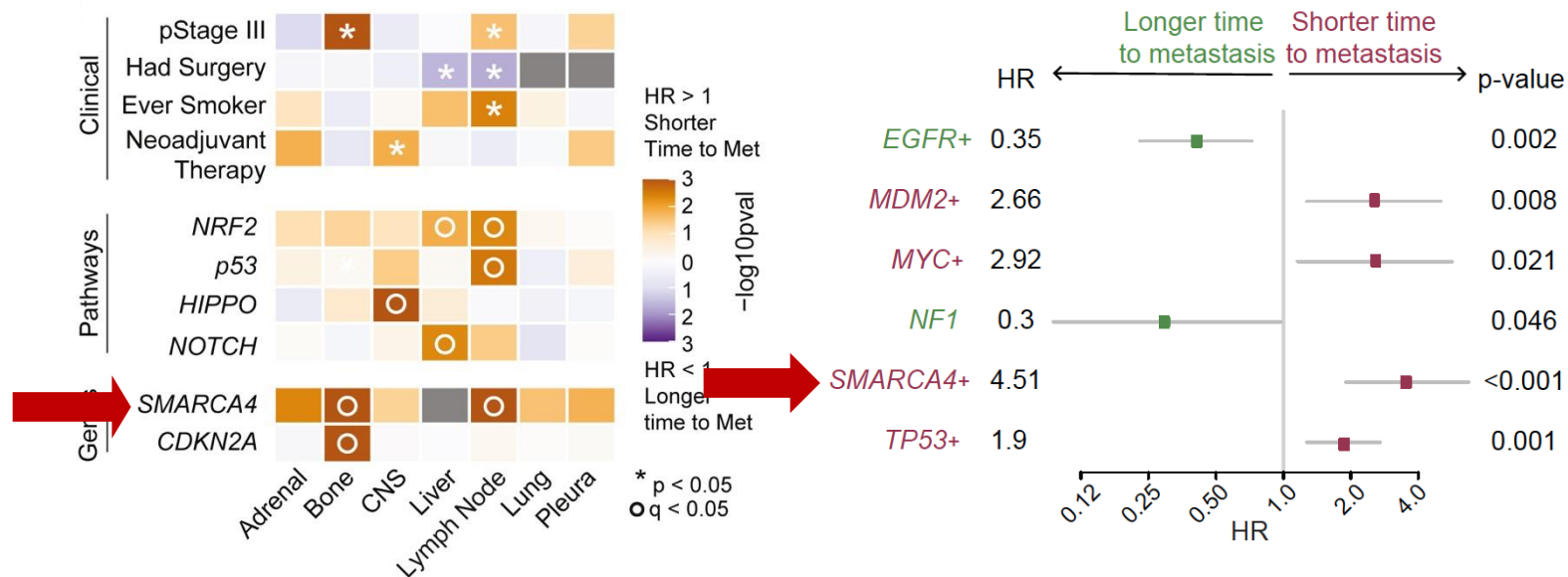
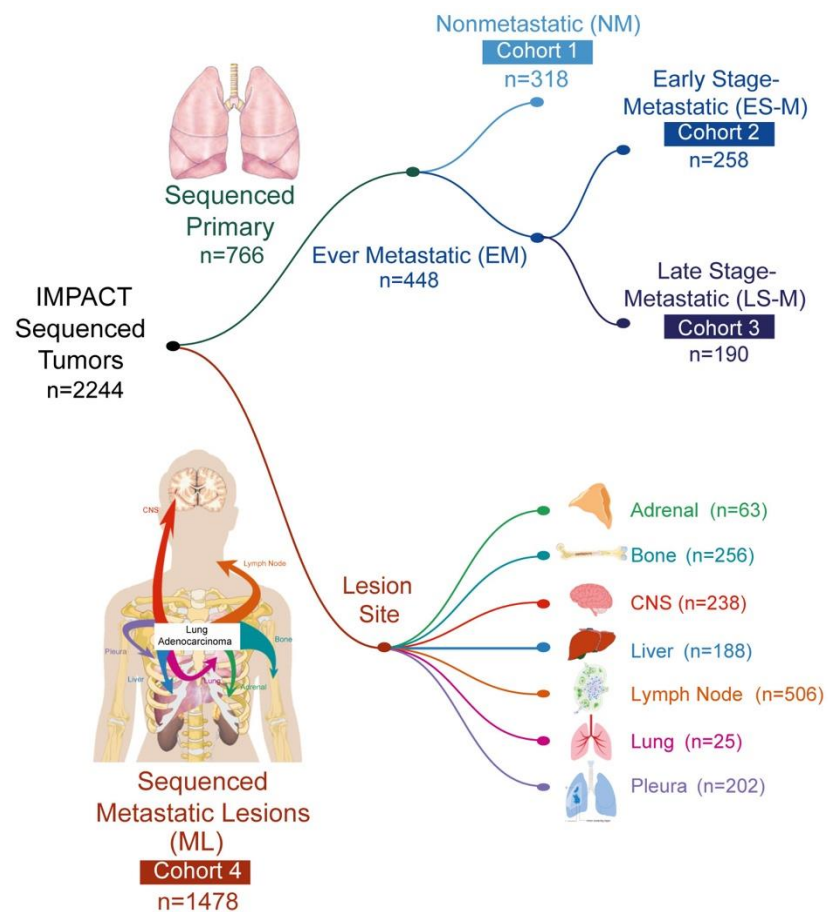
SMARCA4 alterations are associated with poor survival in non-small cell lung cancer



Patients with NSCLC grouped by *SMARCA4* status (n=1288)



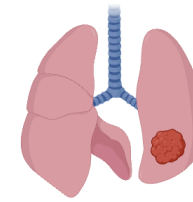
SMARCA4 alterations are associated with shorter time to metastasis



Modeling SMARCA4 deficiency to determine the consequences of chromatin deregulation on tumor evolution



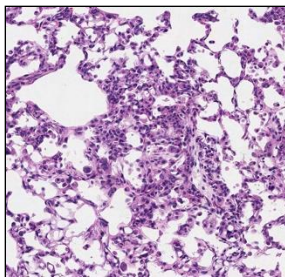
Ad-Cre



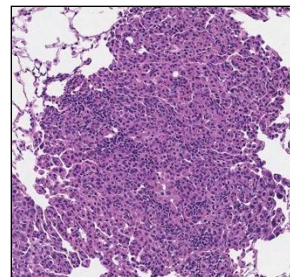
Kras^{LSL-G12D/+}; *Trp53*^{fl/fl} (KP)
Kras^{LSL-G12D/+}; *Trp53*^{fl/fl}; *Smarca4*^{fl/+} (KPS-HET)
Kras^{LSL-G12D/+}; *Trp53*^{fl/fl}; *Smarca4*^{fl/fl} (KPS)

Kras^{G12D/+}; *Trp53*^{-/-} (KP)
Kras^{G12D/+}; *Trp53*^{-/-}; *Smarca4*^{+/-} (KPS-HET)
Kras^{G12D/+}; *Trp53*^{-/-}; *Smarca4*^{-/-} (KPS)

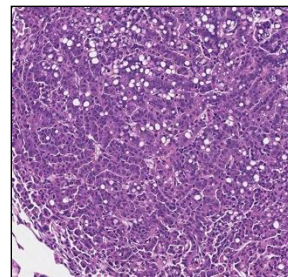
Tumor progression



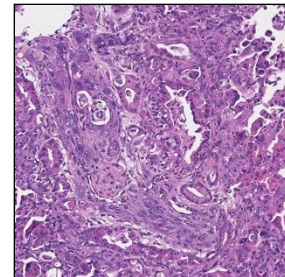
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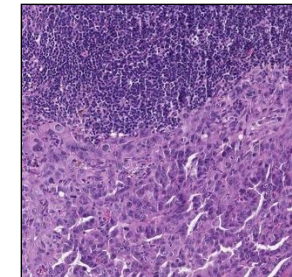
Grade 2



Grade 3

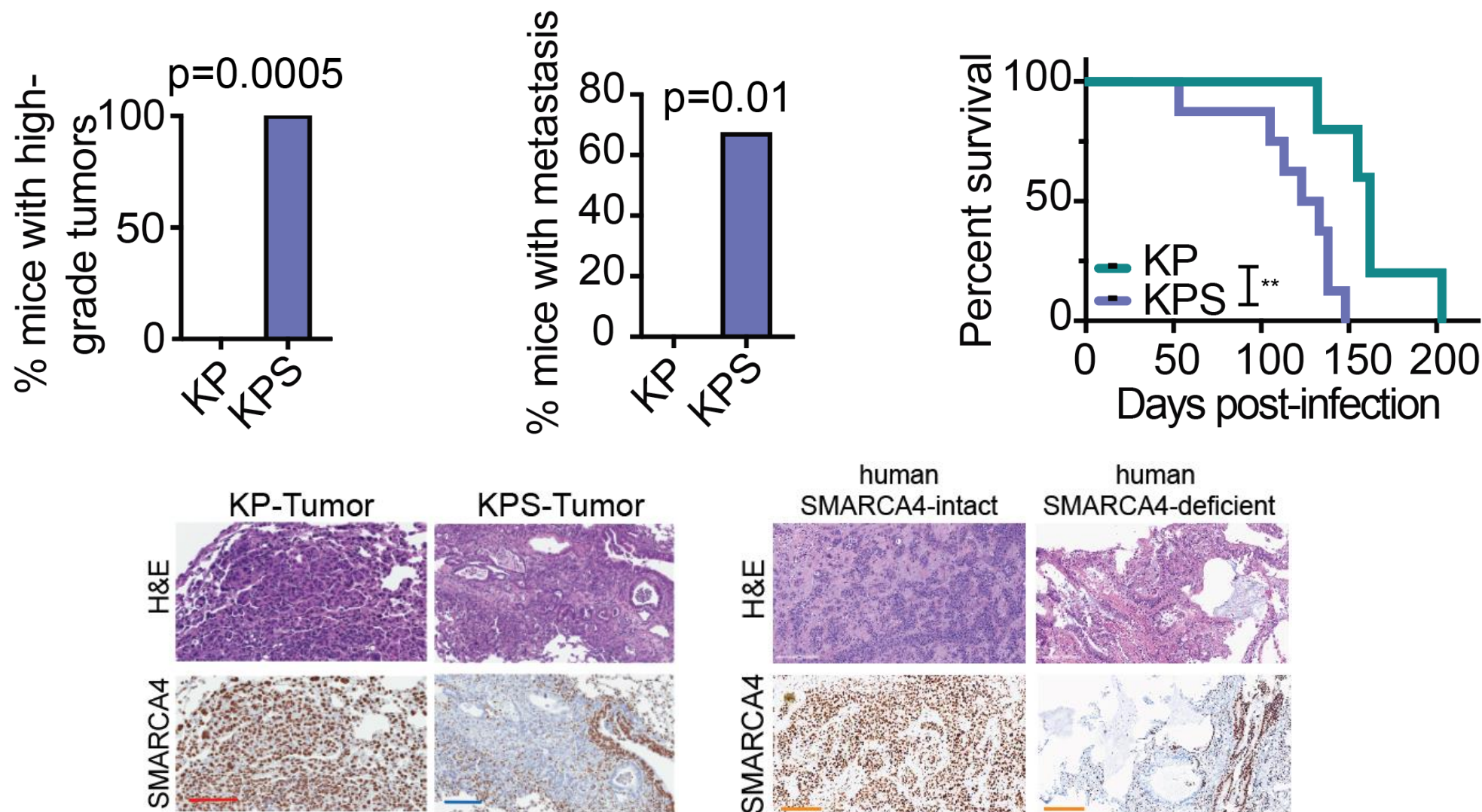


Grade 4

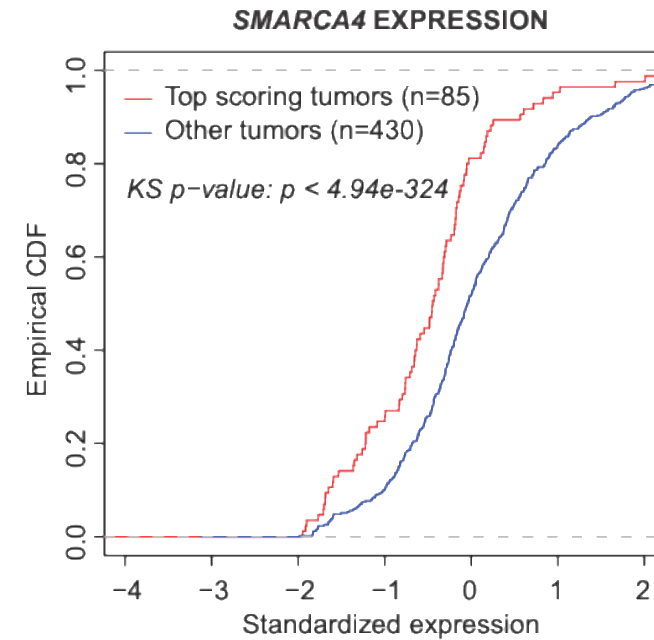
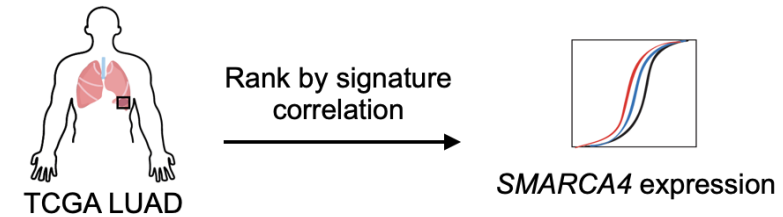
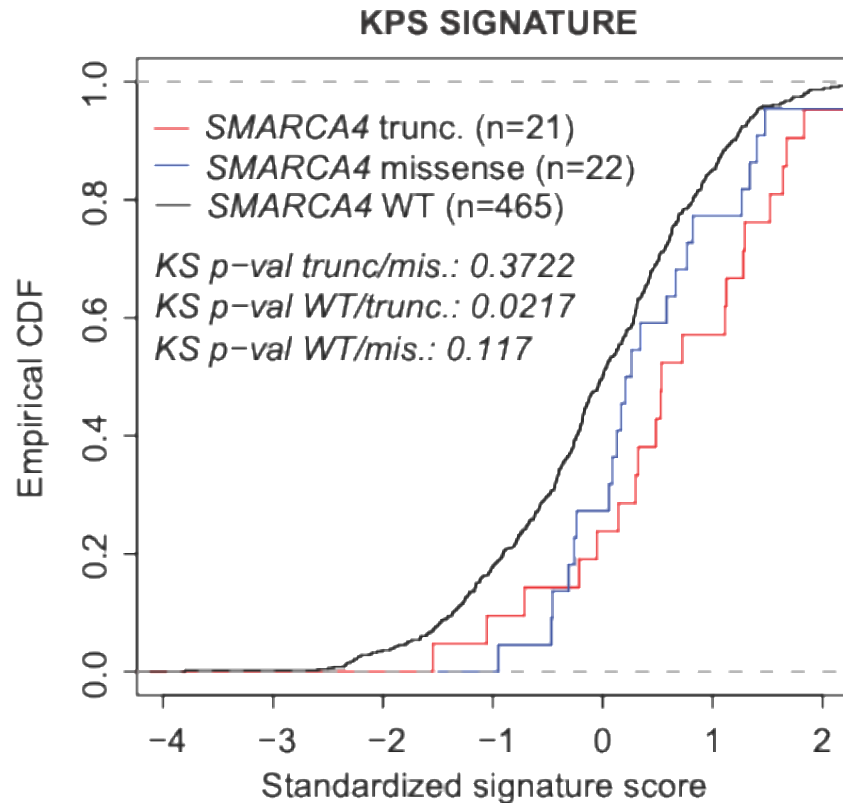


Metastasis

SMARCA4-deficient LUAD GEMMs phenocopy the human disease

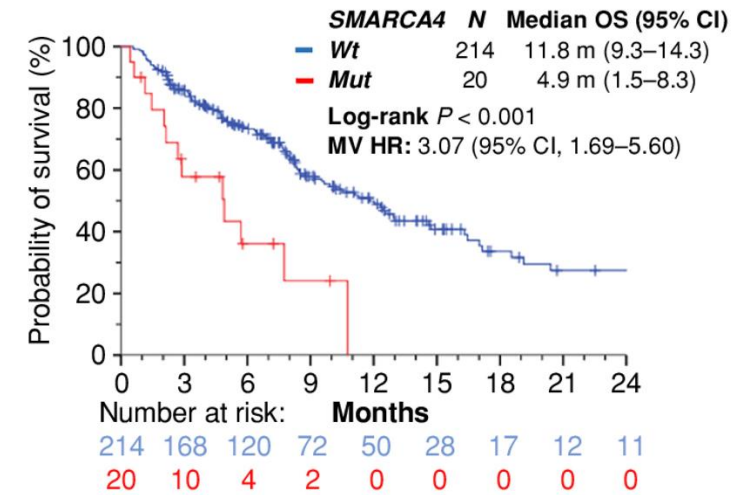
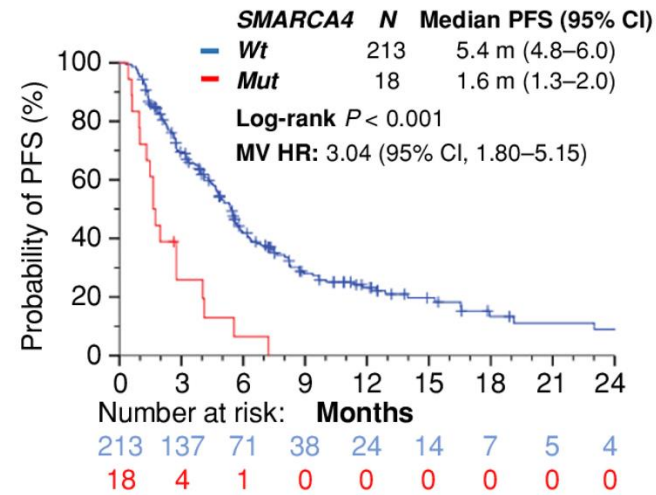
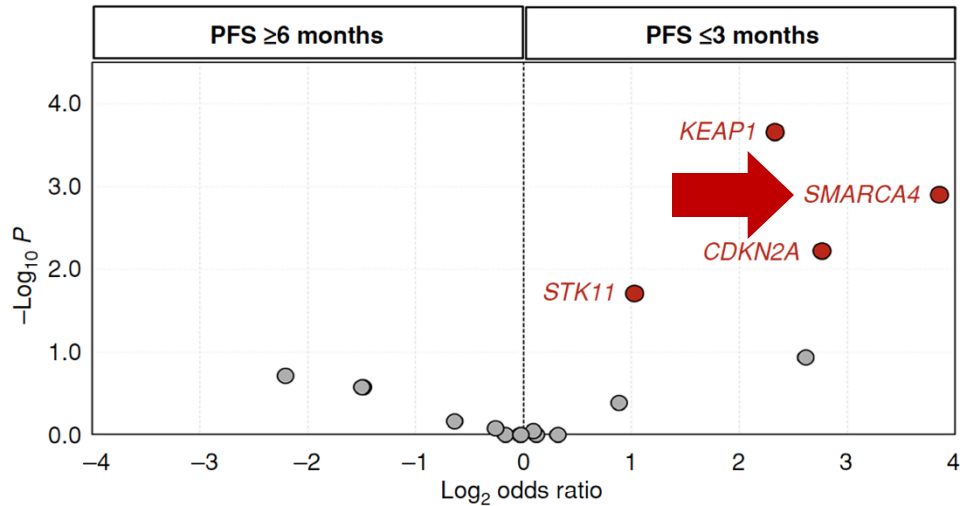


SMARCA4-deficient LUAD GEMMs capture transcriptional states of the human disease

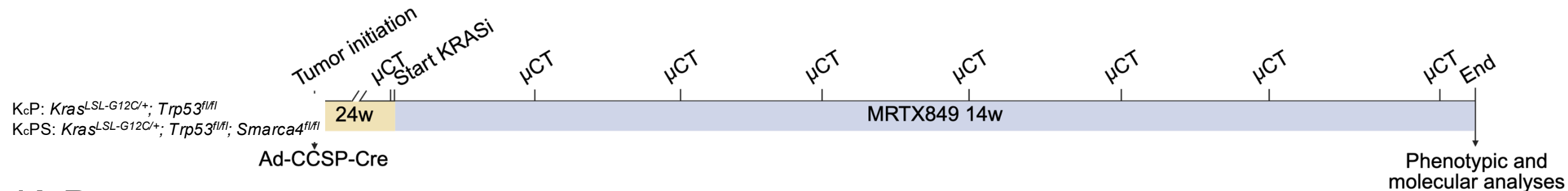


Top scoring tumors ($z > 1$) enriched for *SMARCA4* truncating mutations ($p = 3.18e-3$)

SMARCA4 alterations are associated with poor responses to KRAS-G12C inhibition in the clinic



SMARCA4 deficiency accelerates the onset of acquired resistance



KcP

Pre-treatment

Week 2

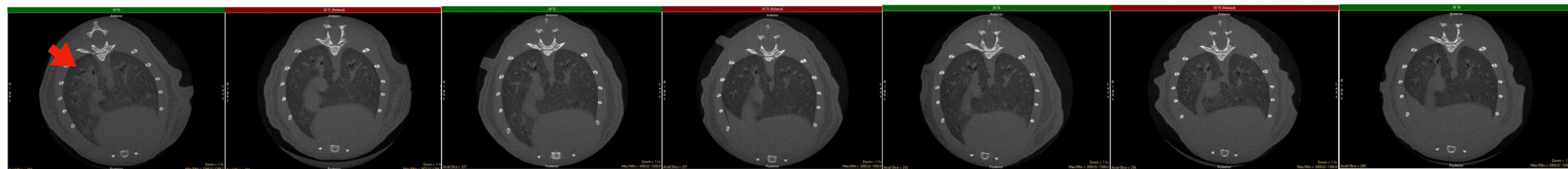
Week 4

Week 6

Week 8

Week 10

Week 12



KcPS

Pre-treatment

Week 2

Week 4

Week 6

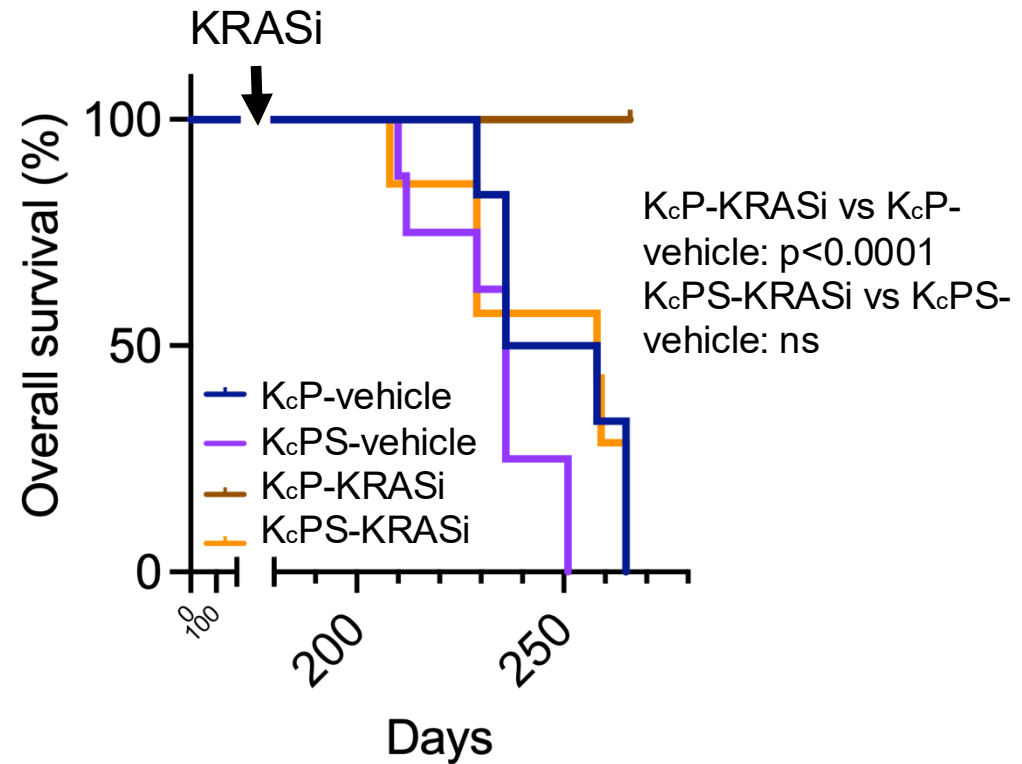
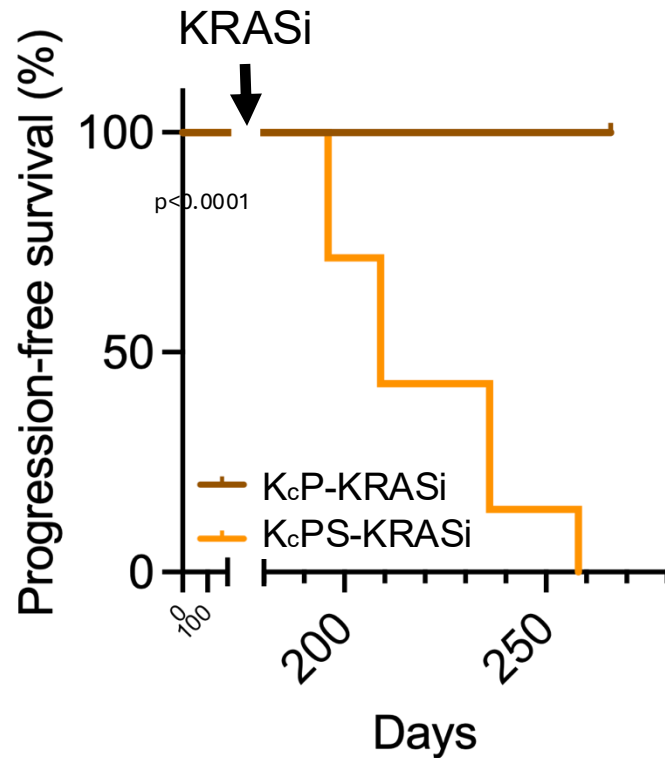
Week 8

Week 10

Week 12



SMARCA4 deficiency abrogates the survival benefit of KRASi

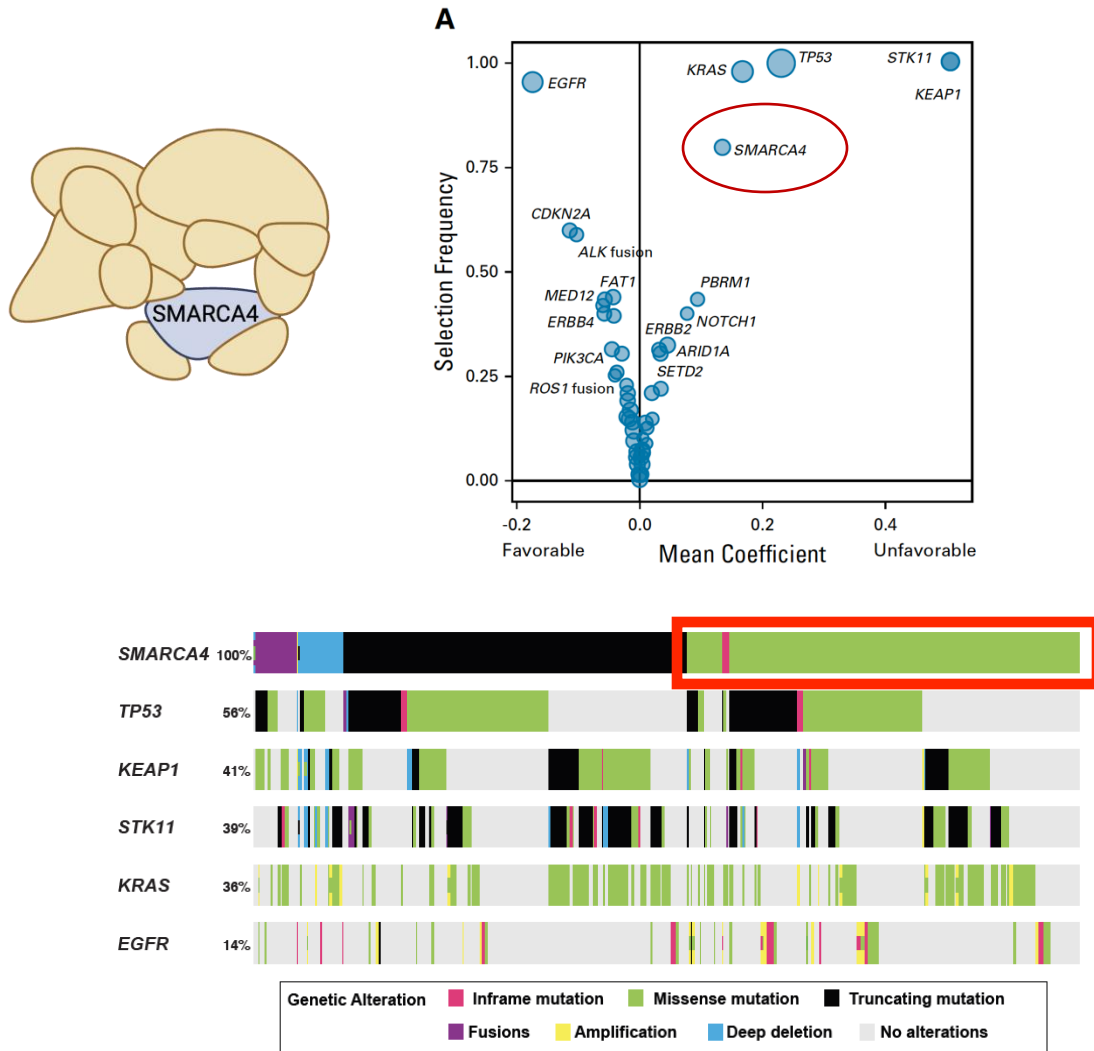


KcP: *Kras*^{LSL-G12C/+}; *Trp53*^{fl/fl}

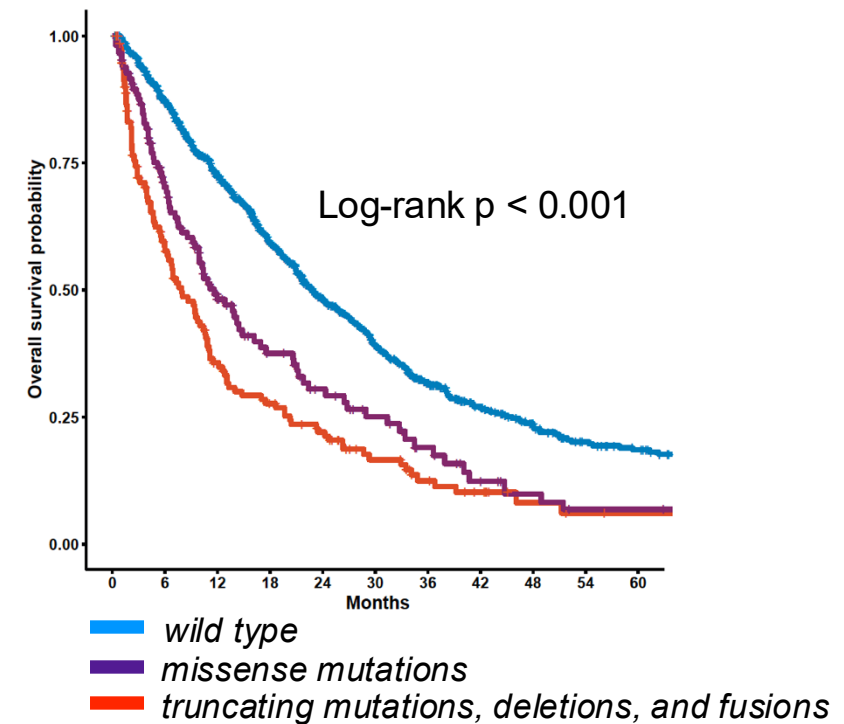
KcPS: *Kras*^{LSL-G12C/+}; *Trp53*^{fl/fl}; *Smarca4*^{fl/fl}

Vicky Liu, unpublished data

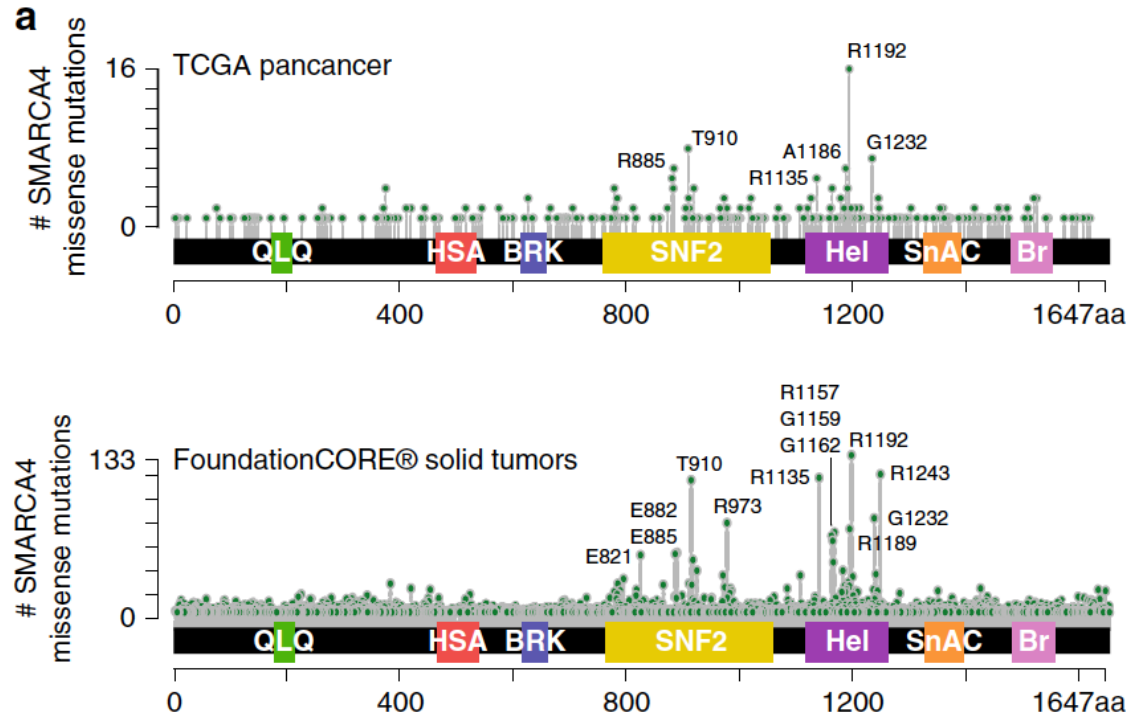
SMARCA4 alterations are associated with poor survival in non-small cell lung cancer



Patients with NSCLC grouped by *SMARCA4* status (n=1288)



Modeling *SMARCA4* cancer variants via prime editing *in vivo*



Lentivirus:



Germline: *Trp53*^{fl/fl}; *Rosa26*^{LSL-PE2}

PMID: 33144586, Zoe Cho

Choose your model **based on the question** you're asking!

Know the **caveats and nuances** of the model you're using or developing. Models are **powerful**. However, a model is “just” a model! **No model is perfect.**

It is critical to **use a variety** of models to fully test a hypothesis.