

A Combined Approach To Treat NSCLC



SAPIENZA
UNIVERSITÀ DI ROMA

•By correction of KRAS G12C through CRISPR-Cas9

•Combined with the novel STAT3 molecular inhibitor W2014-S

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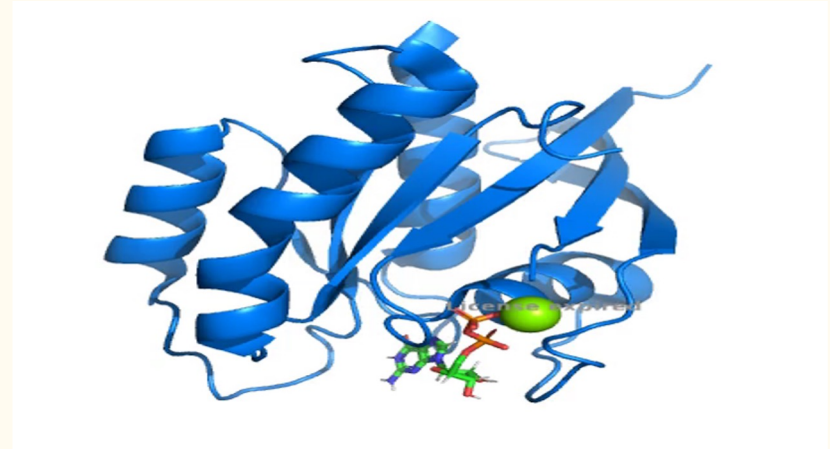
Mirta Mavilia

Background



Non-small cell lung cancer

- Most prevalent type of lung cancer (85% of all cases)
- Frequently diagnosed at an advanced stage, has a poor prognosis.
- 10–25% of patients with adenocarcinoma have KRAS mutation-associated (mostly G12C mutations) tumors



KRAS

- Kirsten RAt Sarcoma virus oncogene homolog from the mammalian ras gene family
- Member of the small GTPase superfamily
- A single amino acid substitution, G12C, causes constant activation of the protein

Background

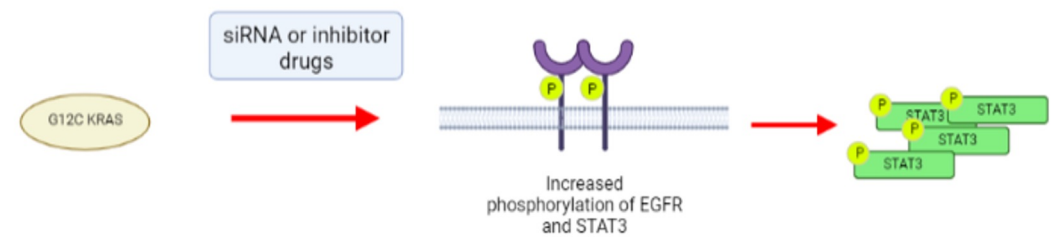
- **EGFR**: regulates epithelial development and homeostasis
- **STAT3**: activated by tyrosine phosphorylation by EGFR

Inhibition of **oncogenic KRAS**, by siRNA or drugs induces phosphorylation and activation of EGFR and STAT3

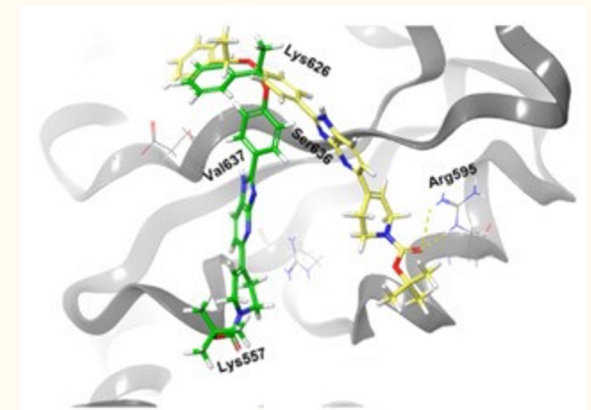
W2014-S, STAT3 Inhibitor

- Derived from Imidazopyridine with one asymmetric center
- Occupies the phosphotyrosine-binding site of STAT3, inhibiting it.

STAT3 & EGFR Correlation with KRAS



Accumulation of EGFR or STAT3 = **tumor formation**



Zheng et al., 2021

Aim of the project

Our strategy: overcoming pEGFR and pSTAT3 accumulation after CRISPR-Cas9 correction

- **Combined therapy** = CRISPR-Cas9 correction for G12C + STAT3 drug inhibitor

Why CRISPR-Cas9?

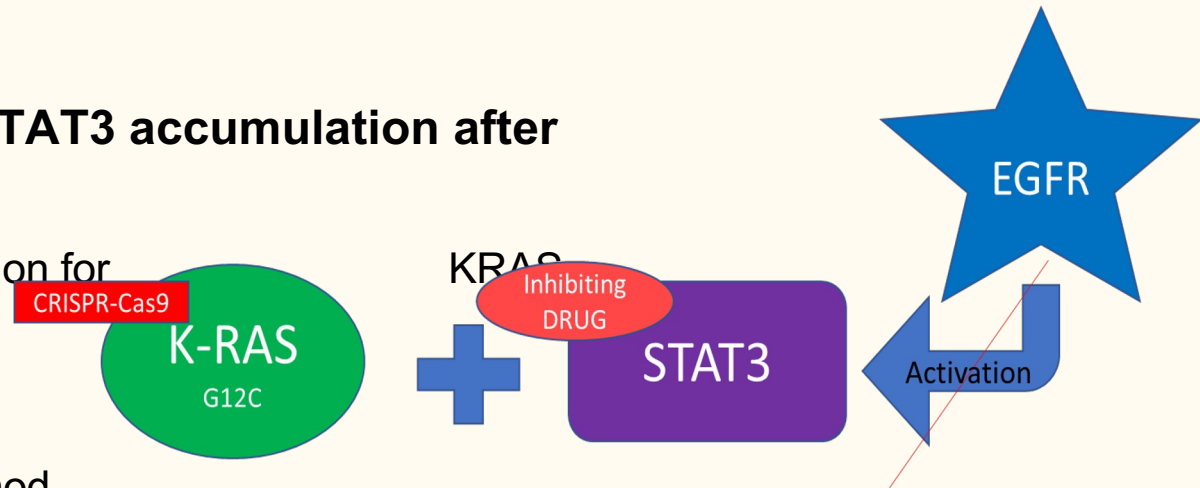
- CRISPR-Cas9 is a long-term correction method
- KRAS G12C inhibition increases pEGFR and pSTAT3 the same is expected after CRISPR-Cas9 G12C correction

Cell line?

- HCC44, human G12C Lung Adenocarcinoma line

Animal model?

- Mouse model: Kras LSL-G12C out mice

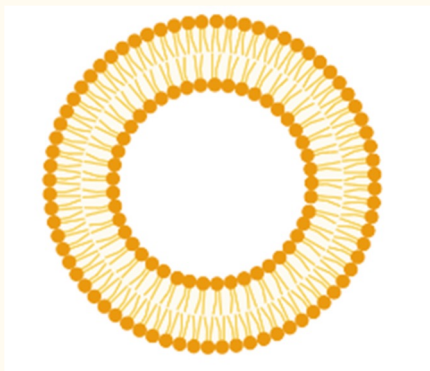


KRAS alleles in the LSL-G12C mouse



Materials and Methods

Delivery method: by liposomes through inhalation

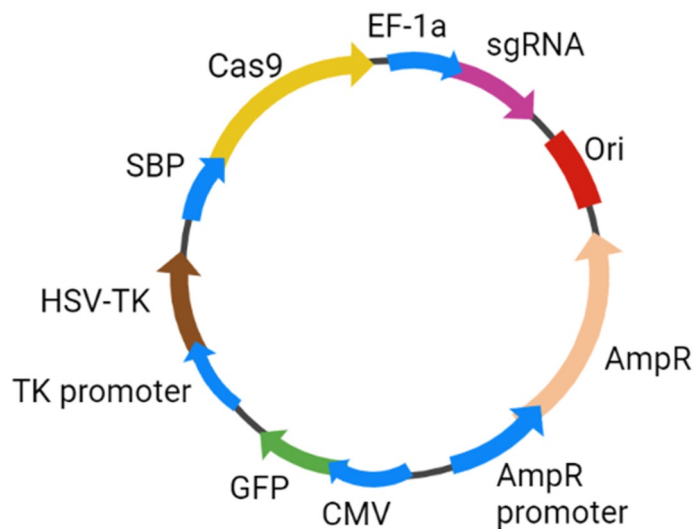


Formulation GL67A:

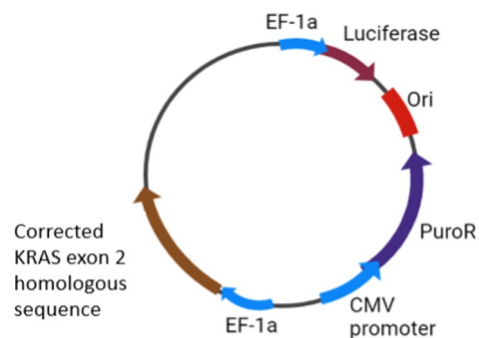
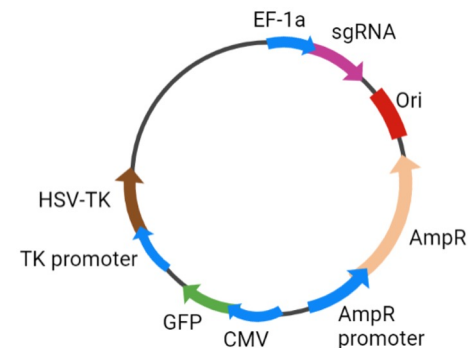
- GL67 cationic lipid
- Zwitterionic colipid DOPE
- Steric stabilizer DP5K

CRISPR-Cas9 system → DSBs

- DSBs repaired with HDR

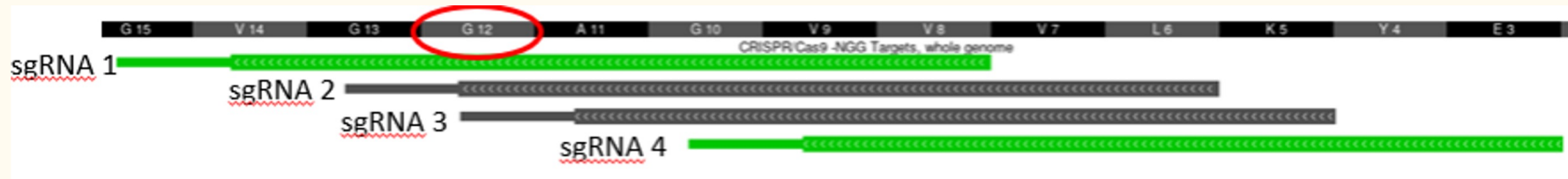


SB-P (surfactant protein B): promoter specific for epithelial lung cells



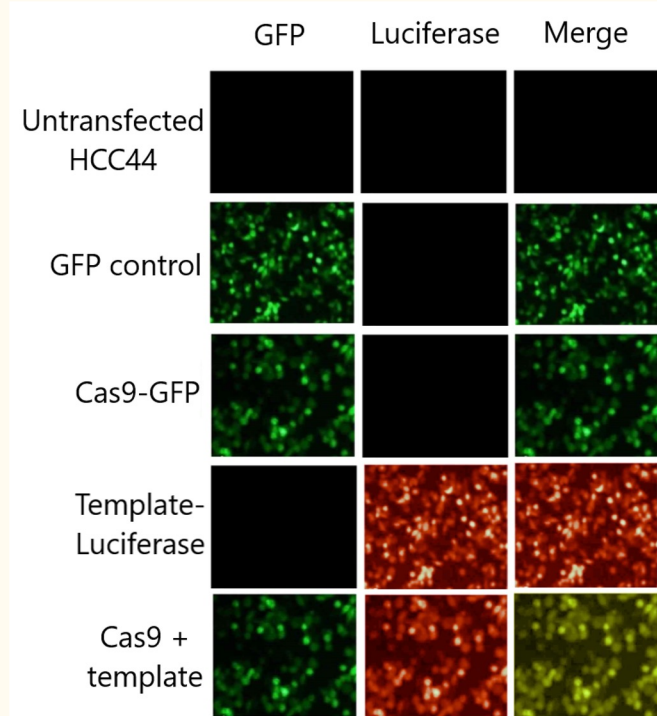
G C C T A C G C C A C C A G C T C C A A C T A C C A C A A G
 G 15 V 14 G 13 G 12 A 11 G 10 V 9 V 8 V 7 L 6

In vitro experiments:



| HCC44 cells | | |
|-------------|-------------------------------|----------------------------|
| | Cleavage at KRAS exon2/ Total | HDR-mediated repair/ Total |
| sgRNA1 | 26/30 | 15/30 |
| sgRNA2 | 3/30 | 1/30 |
| sgRNA3 | 0/30 | 0/30 |
| sgRNA4 | 25/30 | 7/30 |

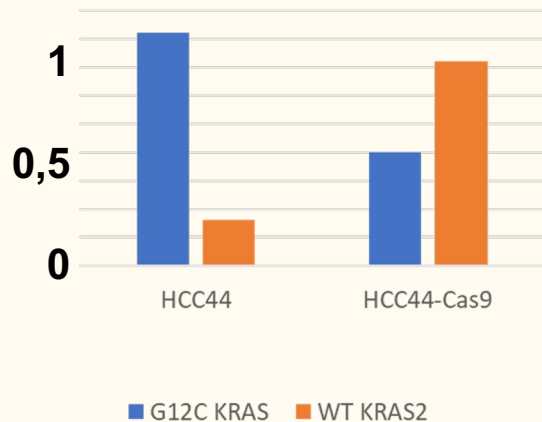
Cleavage and HDR-mediated repair efficiency in different gRNAs



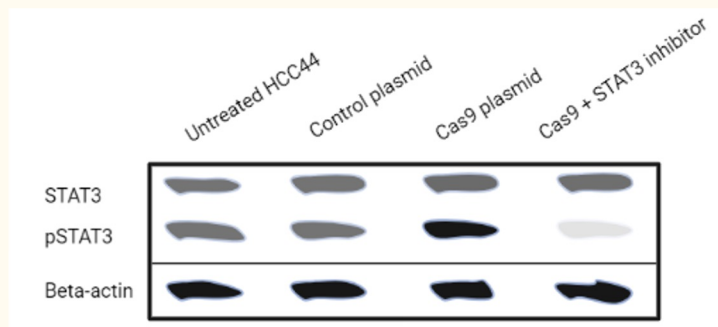
GFP and Luciferase transfection efficiency check

Adapted from <https://genome.ucsc.edu/>, Xu et al., 2017

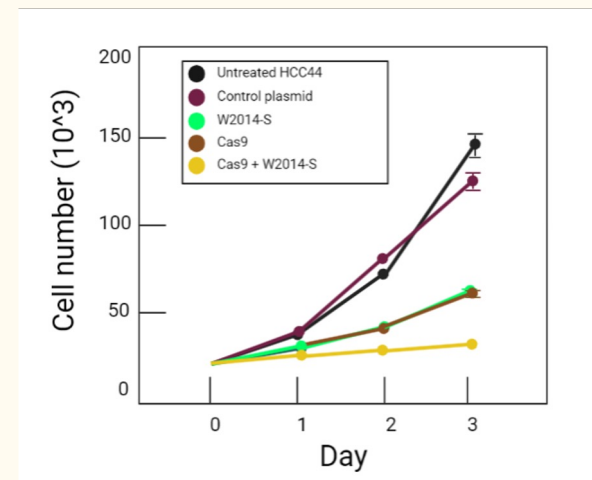
RT-PCR (KRAS/Beta-actin mRNA expression)



pSTAT3 western blot

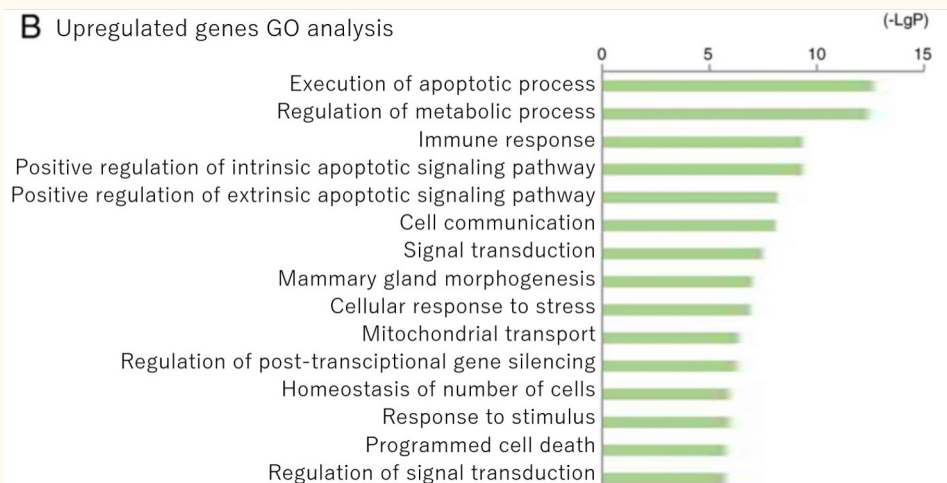


Cell count

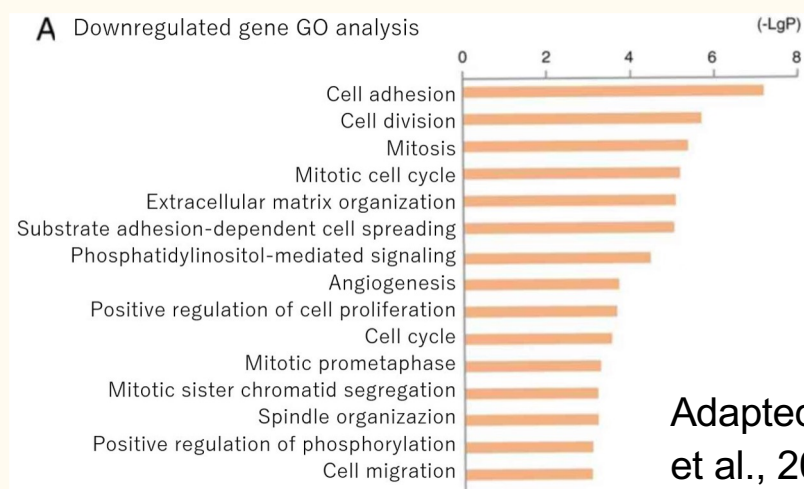


RNA-seq analysis after Cas9 + W2014-S

B Upregulated genes GO analysis

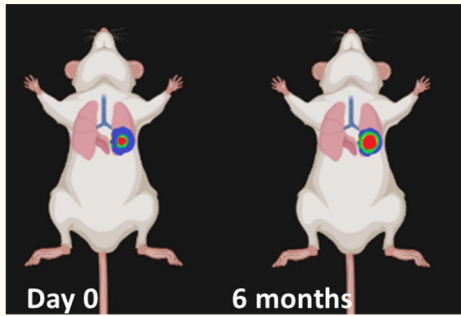


A Downregulated gene GO analysis



Adapted from Zhu et al., 2019

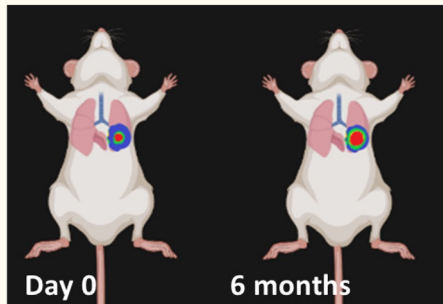
In vivo experiments:



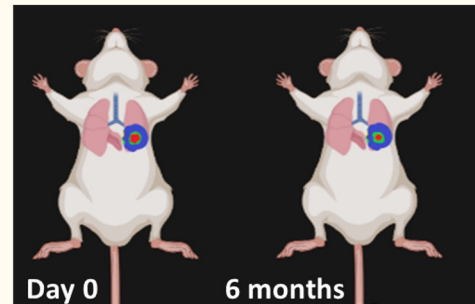
Untreated mouse



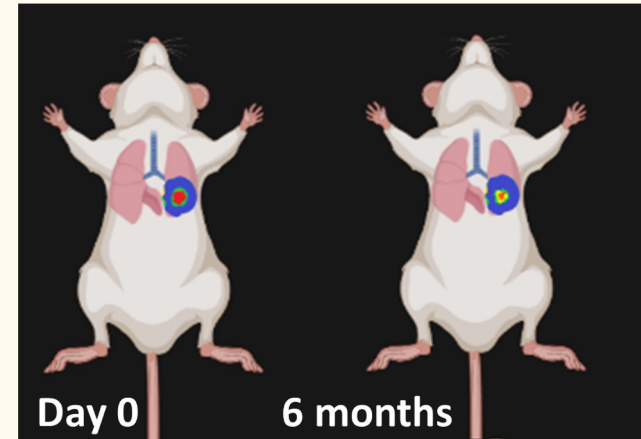
Mouse treated with Cas9 plasmid only



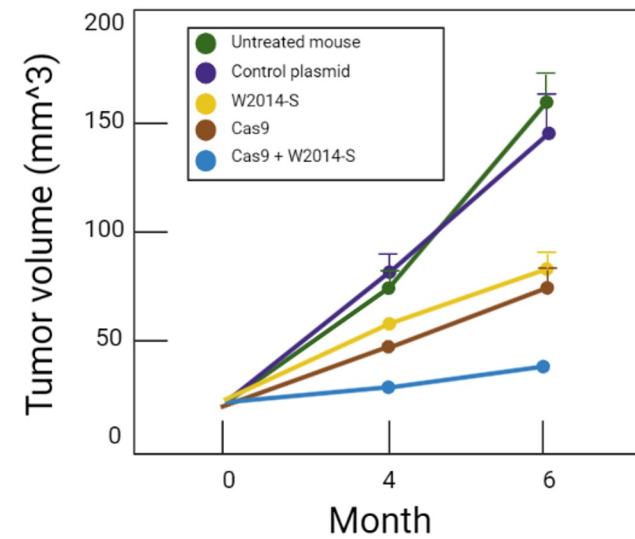
Mouse treated with control plasmid



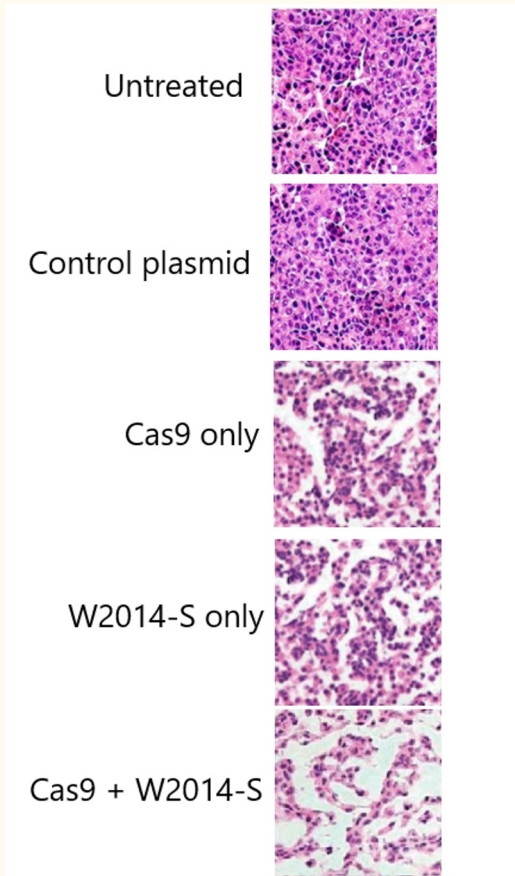
Mouse treated with W2014-S only



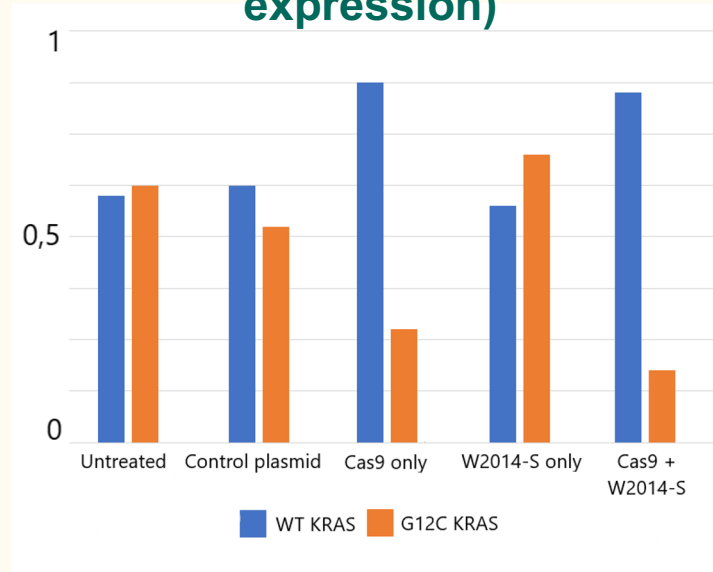
Mouse treated with combination of W2014-S and KRAS G12C correction by CRISPR-Cas9



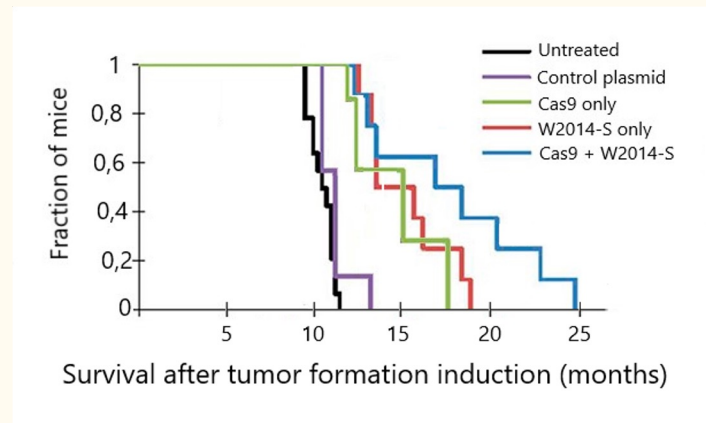
Histopathological analysis



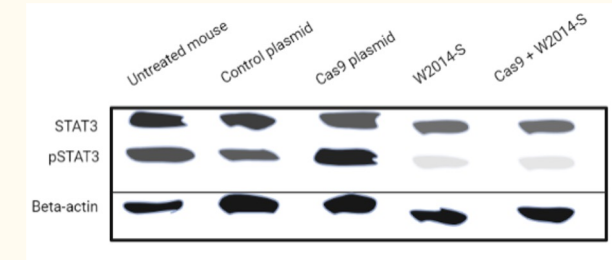
RT-PCR (KRAS/Beta-actin mRNA expression)



Survival curve



Western blot



Guide-seq analysis

| Potential off target sites | Mutated mice |
|------------------------------|--------------|
| intron CYTH4 | 0/12 |
| intron TEX14 | 0/12 |
| intergenic RNF38-MELK | 0/12 |
| exon F11R | 0/12 |
| intron GTF2F1 | 0/12 |
| intergenic CCDC162P-C6orf185 | 0/12 |
| intergenic PCSK6-TM2D3 | 0/12 |
| intergenic LOC401324-HERPUD2 | 1/12 |
| intergenic MIR551A-MEGF6 | 0/12 |

Budget

| Costs/3 years | Budget (€) |
|--------------------------------------------------|--------------------------------|
| Lab equipment | Donated by Sapienza University |
| LSL-KrasG12C Mouse model (excluding maintenance) | 10 000 |
| Researchers' salary (2 Ph students + 1 Post-Doc) | 240 000 |
| Liposome Vector | 8 400 |
| RT-PCR Kit | 600 |
| Western Blot Kit | 400 |
| Crispr-Cas9 Kit | 1 280 |
| HCC44 Cell Line | 400 |
| RNA-Seq | 3 500 |
| GUIDE-Seq | 4 000 |
| Transfection Reagents | 280 |
| Total | 268 860 |

Pitfalls and Solutions

Effects of long-term administration of W2014-S in CRISPR-Cas9 corrected cells haven't been studied yet.



Regular monitoring of STAT3 levels and mice health conditions is necessary.

CRISPR-Cas9 can cause off-target cleavage, leading to mutations.



A modified and more efficient Cas9, like the Sniper-Cas9, could be used instead.

Liposome delivery efficiency is lower than other vectors.



We can add surface molecules like IL4RPep-1 to increase the targeting to lung cancer cells.

Timeline

1st year

Preparation of vectors
In vitro experiments
Data collection

2nd year

In vivo experiments
In vitro replicates
Data collection

3rd year

In vivo replicates
Post-mortem analysis
Survival rate analysis

References

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