# RACGAP1 competitive inhibition in hepatocellular carcinoma via Onyx-015 mRNA transduction

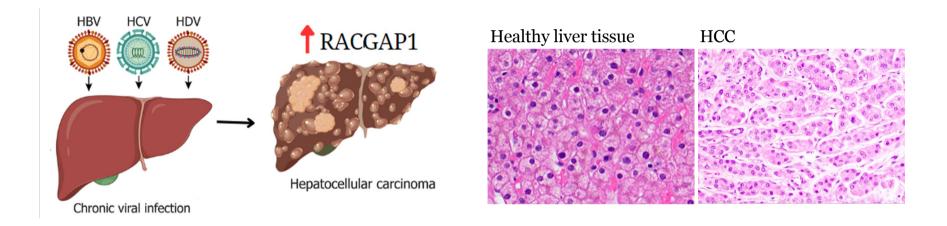




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# Background: Hepatocellular Carcinoma (HCC)

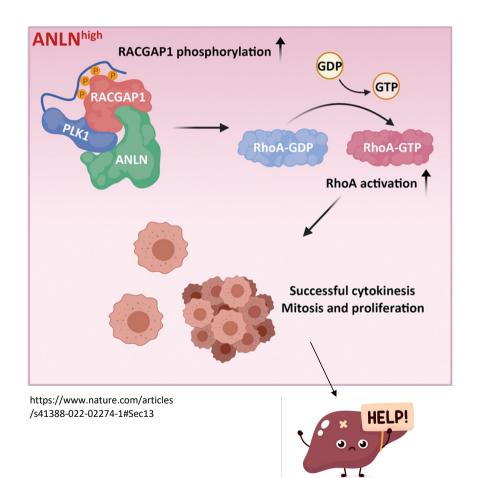


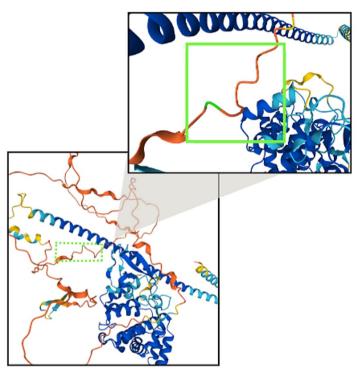
• Liver cancer is the **third most lethal cancer** globally. **Infection by hepatitis B\C** viruses is the main risk factor for HCC development

• The median age: > **60** years

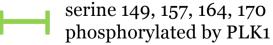
• HCC **recurrence** is significantly associated with **RACGAP1 upregulation**: activation of RACGAP1/Rho/ERK signaling axis

## Background: RACGAP1 pathway





RACGAP1 3D structure via Uniprot



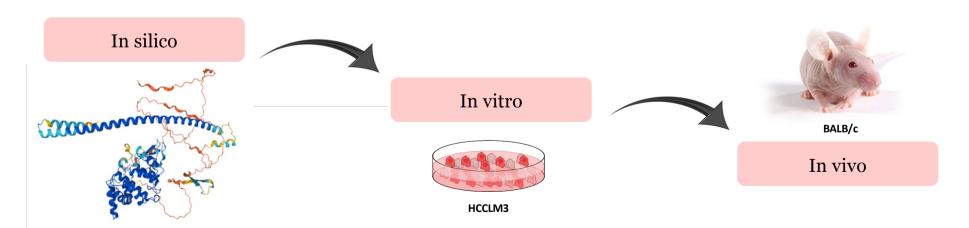
## Aim of the project

### • Induce a **competitive inhibition of RACGAP1** by mutating its phosphorylation sites

• Reduced activation of Rho-A

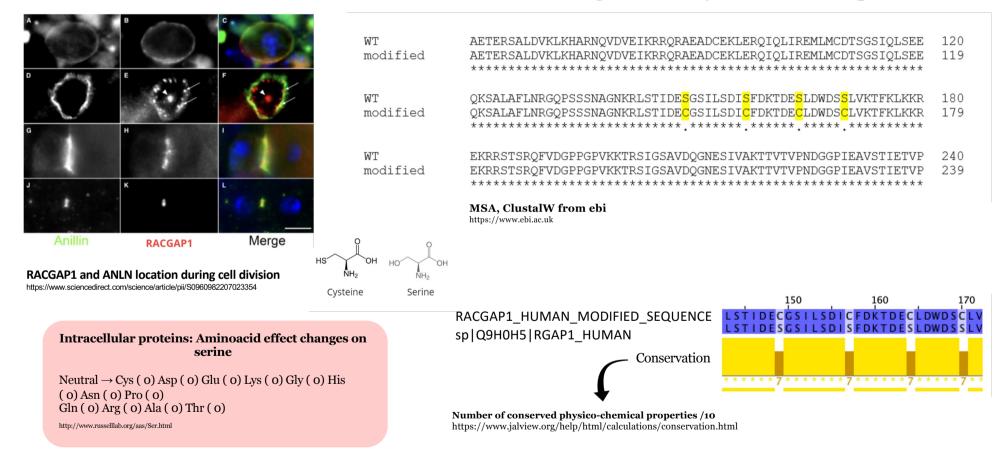
• Inhibition of self proliferation and decrease in HCC size

### How do we do it?



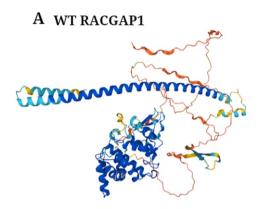
## In silico

### Amino acid modifications: does the aminoacidic change cause any effects on the protein?



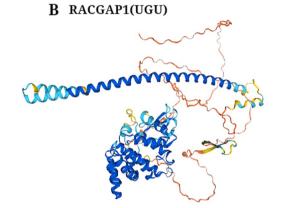
## In silico

### Structural predictions



Structure prediction via alphaphold

https://www.uniprot.org/uniprotkb/Q9H0H5/featur e-viewer



Structure prediction via Swissprot

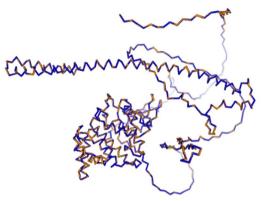
https://swissmodel.expasy.org/interactive/Xk9YBQ /models/

#### Model Confidence:

Very high (pLDDT > 90) Confident (90 > pLDDT > 70) Low (70 > pLDDT > 50) Very low (pLDDT < 50)

https://www.uniprot.org/uni protkb/Q9H0H5/featureviewer

#### C WT RACGAP1/RACGAP1(UGU)

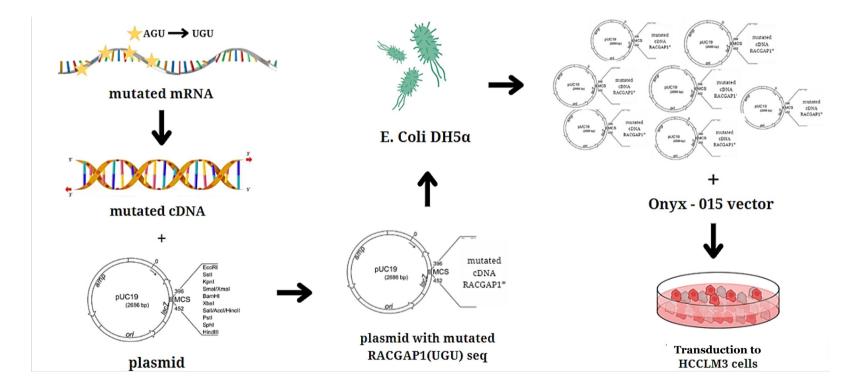


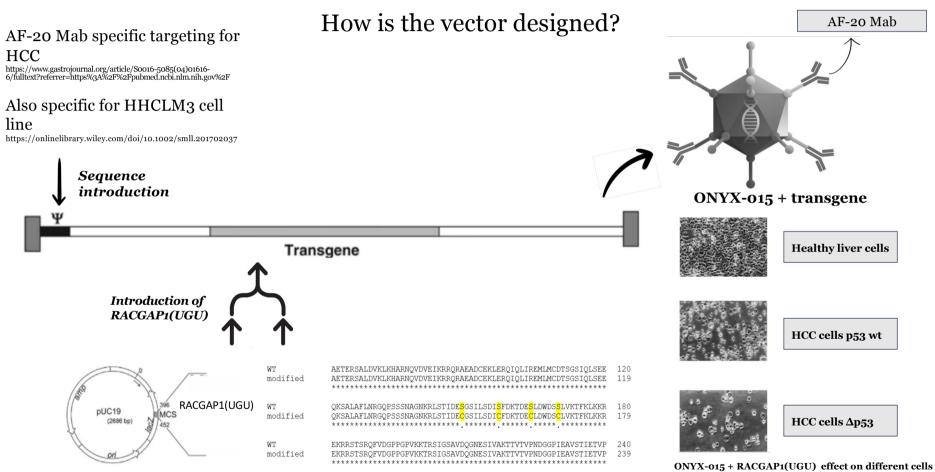
3D structure superposition via DALI

Legend: **Structure conservation Dark blue regions are structurally aligned** http://ekhidna2.biocenter.helsinki.fi/dali/DaliTuto rial.pdf

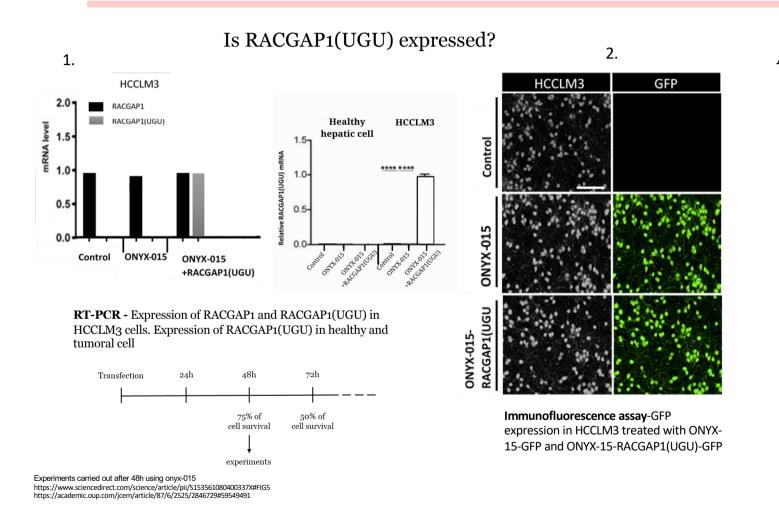
Z-score=46.8 Significant similarities' have a Z-score above 2 https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2 639270/

### Cloning and transduction of RACGAP1(UGU) mRNA

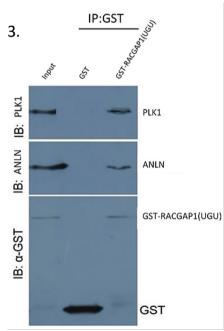




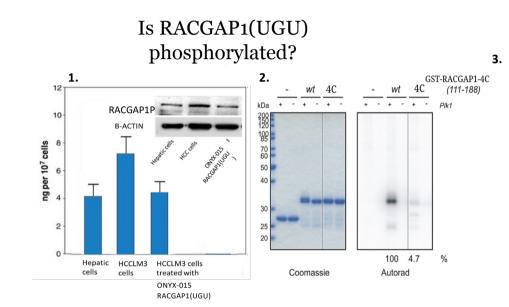
https://images.app.goo.gl/86RRRzDvhZAAyWtd6



#### Does the RACGAP1(UGU)-ANLN-PLK1 complex form?

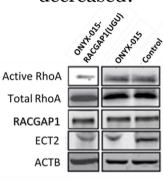


**Pull down-** RACGAP1(UGU)-ANLN-PLK1 complex formation



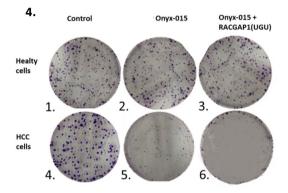
**Phosphorylation assay** - ELISA and Western Blot - Normal levels of RACGAP1P in healthy cells, elevetated levels of RACGAP1P in HCCLM3, reduced levels of RACGAP1P in treated HCCLM3 GST(-), GST-RACGAP1-111/188 (Wt), and GST-RACGAP1-4C-111/188 (4C) were incubated with Plk1 and [ $\gamma$ -32P] ATP. Relative incorporation of 32P to GST-RACGAP1 was quantified. (111-188=peptide from amino acid 111 to 188)

# Is RohA activity decreased?



Western blot - Detection of RhoA activity and also ECT2 and RACGAP1 expression in HCCLM3 after trasfection of RACGAP1(UGU) Adapted from (Yang et al., 2018)

# What happens to the cells?



#### **Clonogenic assay:**

1. Healthy Hepatic cells,

2. Healthy Hepatic cells with transduction of empty Onyx-015,

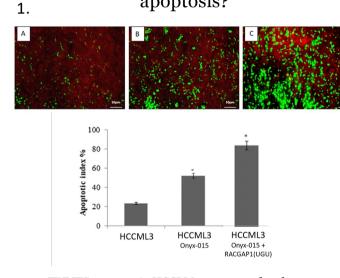
3. Healthy Hepatic cells with transduction of RACGAP1(UGU) mutated protein,

4. Hepatocarcinoma HCCLM3 cells,

5. Hepatocarcinoma HCCLM3 with transduction of empty Onyx-015,

6. Hepatocarcinoma HCCLM3 with transduction of RACGAP1(UGU) mutated protein,

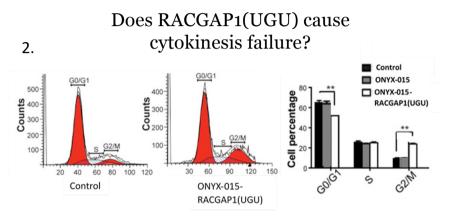
#### Does RACGAP1(UGU) cause apoptosis?



**TUNEL assay - A.** HCCLM3 non treated and no apoptotic cells are detected. **B.** HCCLM3 treated with emptyvector, no apoptotic cells are detected **C.** HCCLM3 treated with the mutated RACGAP1(UGU), incresead levels of apoptotic cells

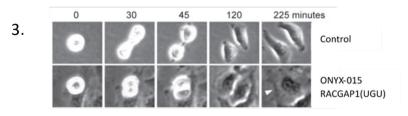
#### Adapted from

https://www.researchgate.net/publication/335679404\_In\_Vivo\_Anti-Tumor\_Effects\_of\_Citral\_on\_4T1\_Breast\_Cancer\_Cells\_via\_Induction \_of\_Apoptosis\_and\_Downregulation\_of\_Aldehyde\_Dehydrogenase\_Ac tivity



**Cell cycle analysis - 1.** Cell count in different cell cycle phases, RACGAP1(UGU) vs control HCCLM3 cells.

**2.** Cell percentage in different cell cycle pahses, RACGAP1(UGU) vs control HCCLM3 cells.

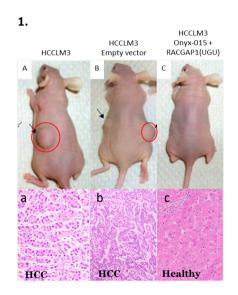


**Cytokinesis analysis** - Selected frames from time-lapse imaging of ONYX-015-RACGAP1(UGU) and control HCCLM3 cells

Adapted from Adapted from (Yang et al., 2018)

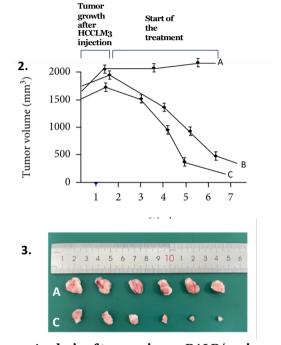
## In vivo

### Is there a change in the tumor mass?



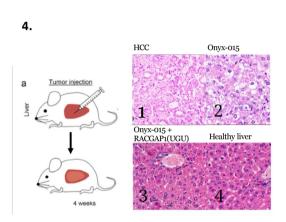
**Tumor detection** in BALB/c mouse injected with **A.** HCCLM3 cells; **B**. HCCLM3 cells + empty vector; and **C.** HCCLM3 cells + onyx-015 + RACGAP(UGU) and **relative histological samples.** 

Adapted from https://bmccancer.biomedcentral.com/articles/10.118 6/1471-2407-11-425/figures/7



2.Analysis of tumor size on BALB/c mice injected with
A. HCCLM3
B. HCCLM3 treated with empty vector
C. HCCLM3 treated with onyx-015 +
RACGAP(UGU)

**3.Representative images of tumors** removed from BALB/c mice from samples A and C



### HCC histological samples from C57BL/6 mice

- 1. HCC tissue
- **2.** Tissue sample injected with empty vector **3.** Tissue sample injected with RACGAP1(UGU) vector
- **4.** Healthy liver tissue

Adapted from https://pubmed.ncbi.nlm.nih.gov/15649325/

https://www.hindawi.com/journals/omcl/2022/3034150/

## Project budget

### Cloning and transduction

mRNA  $\rightarrow$  \$5 640 (\$10/RNA base) plasmid  $\rightarrow$  \$94 vector for in vitro and vivo  $\rightarrow$  \$2670 sequencing  $\rightarrow$  \$75 **Total**  $\rightarrow$  **\$8479** 

### In vitro

- reverse transcription  $\rightarrow$  \$490
- cell line  $\rightarrow$  \$700
- plasmids  $\rightarrow$  \$800
- $qPCR \rightarrow \$800$
- $co-ChIP \rightarrow $700$

- Elisa  $\rightarrow$  \$700
- Western blot  $\rightarrow$  \$700
- clonogenic assay  $\rightarrow$  \$200
- TUNEL  $\rightarrow$  \$600
- cell cycle analysis  $\rightarrow$  \$350
- Total→ \$6040

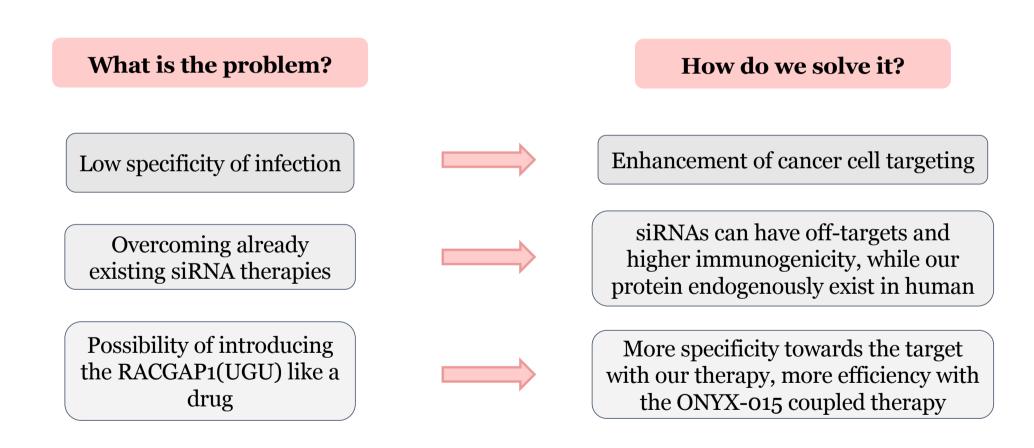
#### In vivo

4 BALB/c nude mouse per group x3 (36 tot)  $\rightarrow$  \$2160 4 C57BL/6 mouse x3 (36 tot) $\rightarrow$  \$720 Mice maintenance  $\rightarrow$  \$10 000 **Total**  $\rightarrow$  **\$2880** 

#### Salaries

3 PhD  $\rightarrow$  \$60 000/year 2 post DOC  $\rightarrow$  \$ 50 000/year Total  $\rightarrow$  \$110 000/year Total for three years of work \$357 399

## Pitfalls and solutions



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