

# Targeting Pin1 in cholangiocarcinoma tumor and stromal cells using siRNA therapeutics

Justin H. Lo,<sup>1,2</sup> Thatcher R. Heumann,<sup>1</sup> Nora Francini,<sup>2</sup> Eva F. Gbur,<sup>2</sup> Michelle Zhou,<sup>2</sup> Shabnam Z. Eghbali,<sup>3</sup> Laura W. Goff,<sup>1</sup> and Craig L. Duval<sup>2</sup>

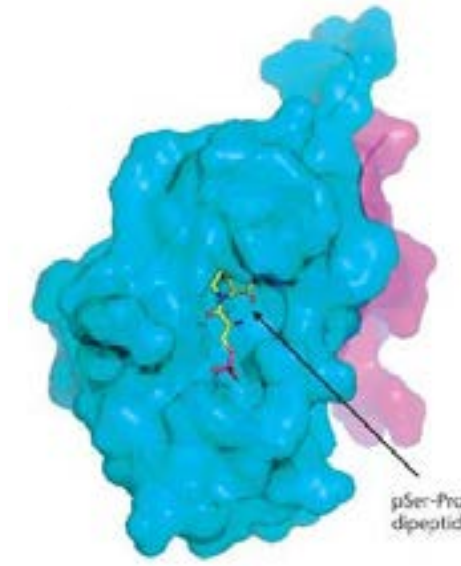
<sup>1</sup>Division of Hematology/Oncology, Dept. of Medicine, Vanderbilt University Medical Center; <sup>2</sup>Department of Biomedical Engineering, Vanderbilt University;

<sup>3</sup>Department of Medicine, Vanderbilt University Medical Center

## Introduction

### Pin1 as a therapeutic target

- Pin1 (peptidyl-prolyl cis-trans isomerase NIMA-interacting 1) overexpression is seen in many tumor types and has been correlated with poor prognosis<sup>1</sup>
- Previous studies in pancreatic cancer have associated higher Pin1 expression with an immunosuppressive tumor microenvironment (TME)<sup>2</sup>
- Inhibition of Pin1 in animal models renders pancreatic cancer eradicable by chemoimmunotherapy<sup>2-3</sup>
- Like pancreatic cancer, cholangiocarcinoma (CCA) carries a poor prognosis and is characterized by desmoplastic stroma<sup>4</sup> *Pin1 structure*<sup>5</sup>
- We hypothesized that targeting Pin1 in CCA could be a well-tolerated approach with therapeutic benefit, in part by enhancing anti-tumor immune response to CCA

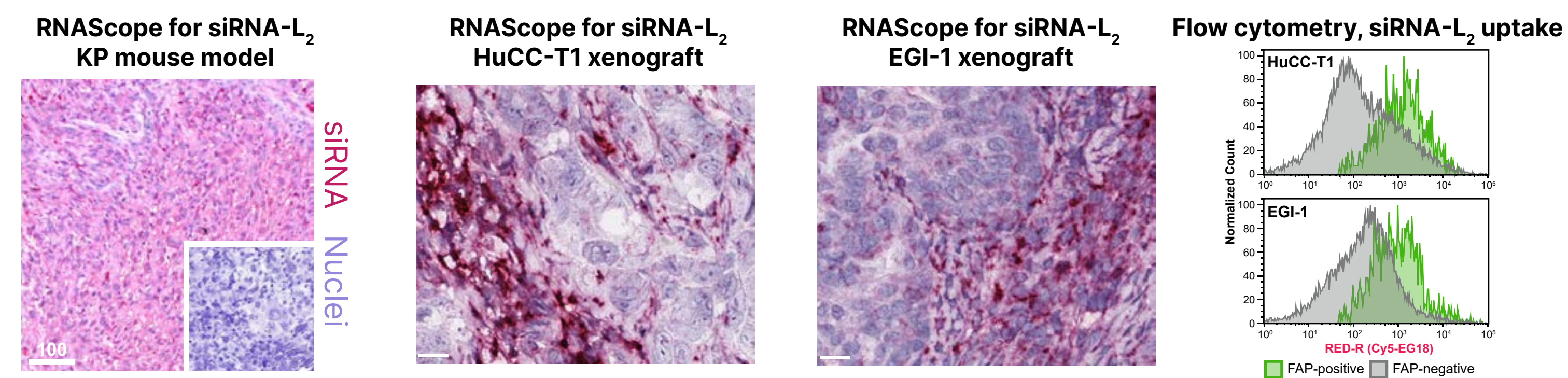


### Rationale for targeting Pin1 using siRNA therapeutics

- Small interfering RNA (siRNA) can be designed to be specific for Pin1 and employ delivery methods that are tumor-tropic
- Other methods of targeting PIN1 exist, but have shortcomings:
  - All-trans retinoic acid (ATRA) +/- arsenic trioxide (ATO) are not molecularly specific for Pin1 and can cause various side effects such as cardiotoxicity or cytopenias (for ATO) and hypertriglyceridemia or teratogenicity (for ATRA)
  - A small molecule Pin1 inhibitor, sulfopin, is also available but does not have tumor tropism/selectivity

## Biodistribution and Safety

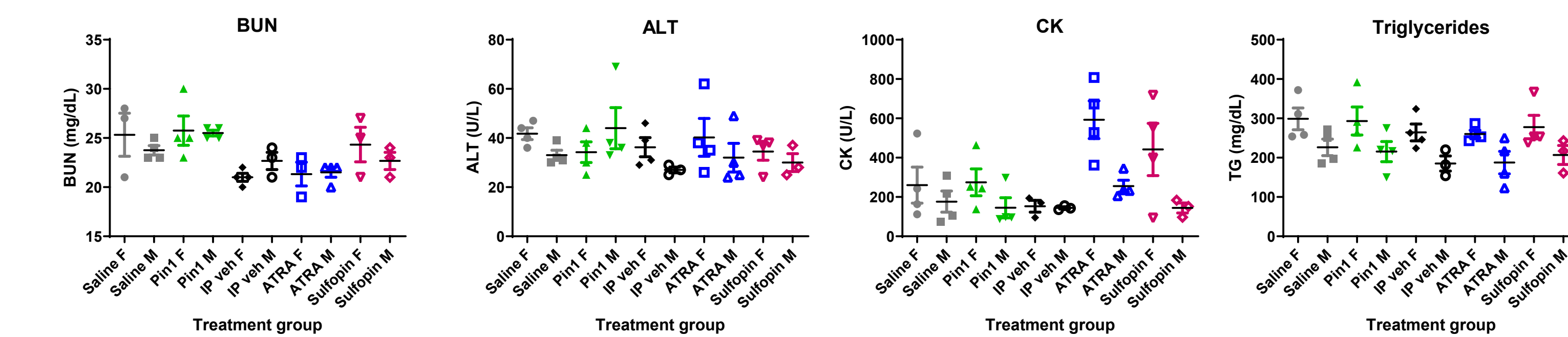
### siRNA-L<sub>2</sub> accumulation in mouse and human tumors



### Safety/tolerability of Pin1 siRNA-L<sub>2</sub> administration in healthy mice

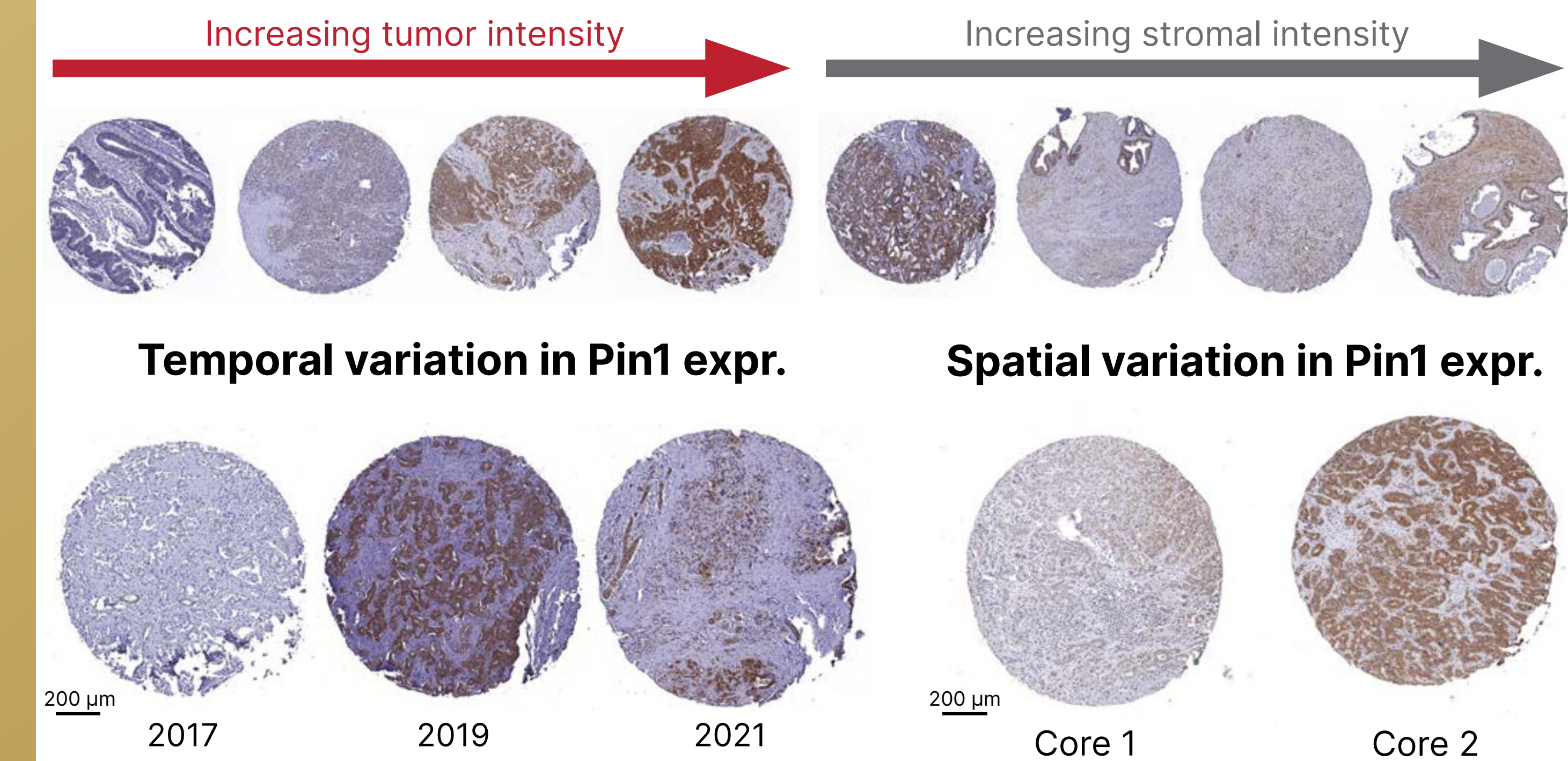
Healthy male and female FVB mice were treated for one week with the following (n=4 per sex per group):

- Pin1 siRNA-L<sub>2</sub>, 20 mg/kg i.v. twice a week
- Saline, i.v. twice a week
- ATRA, 10 mg/kg i.p. daily
- Sulfopin, 40 mg/kg i.p. daily
- Vehicle only, i.p. daily

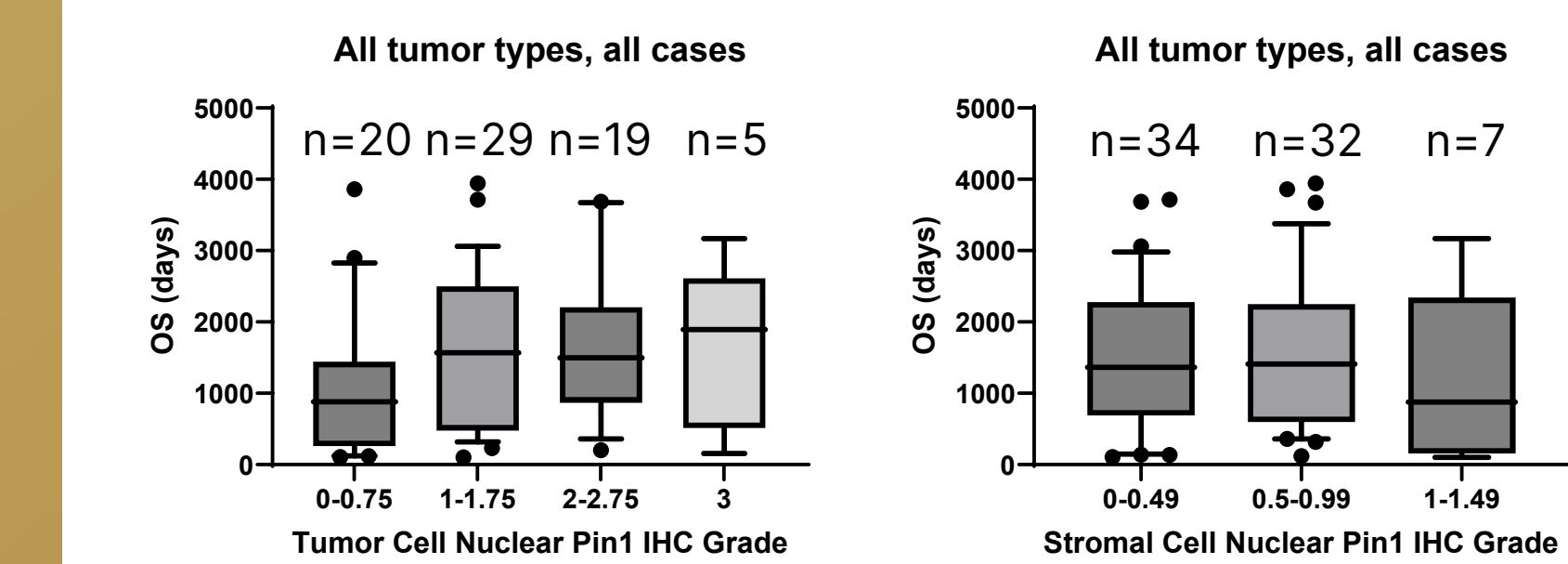


## Pin1 in Human CCA Tumor Microarray

### Examples of Staining Patterns



### Pin1 intensity vs. overall survival

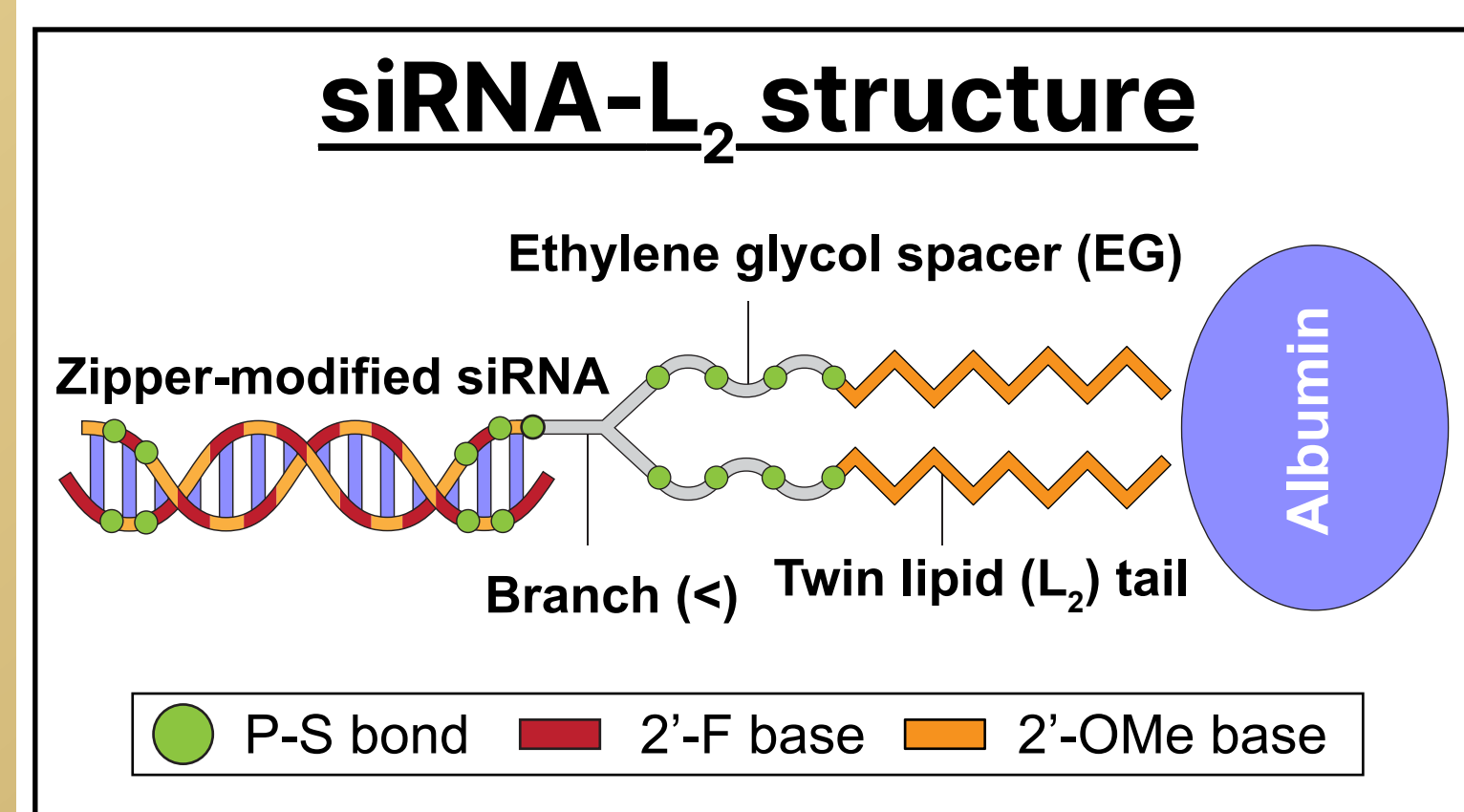


Analyses by sex, tumor stage/resectability, and tumor location (intrahepatic vs. extrahepatic) are in process, as well as different outcome measures including cancer recurrence and treatment response. Grading by cytoplasmic rather than nuclear expression is also underway.

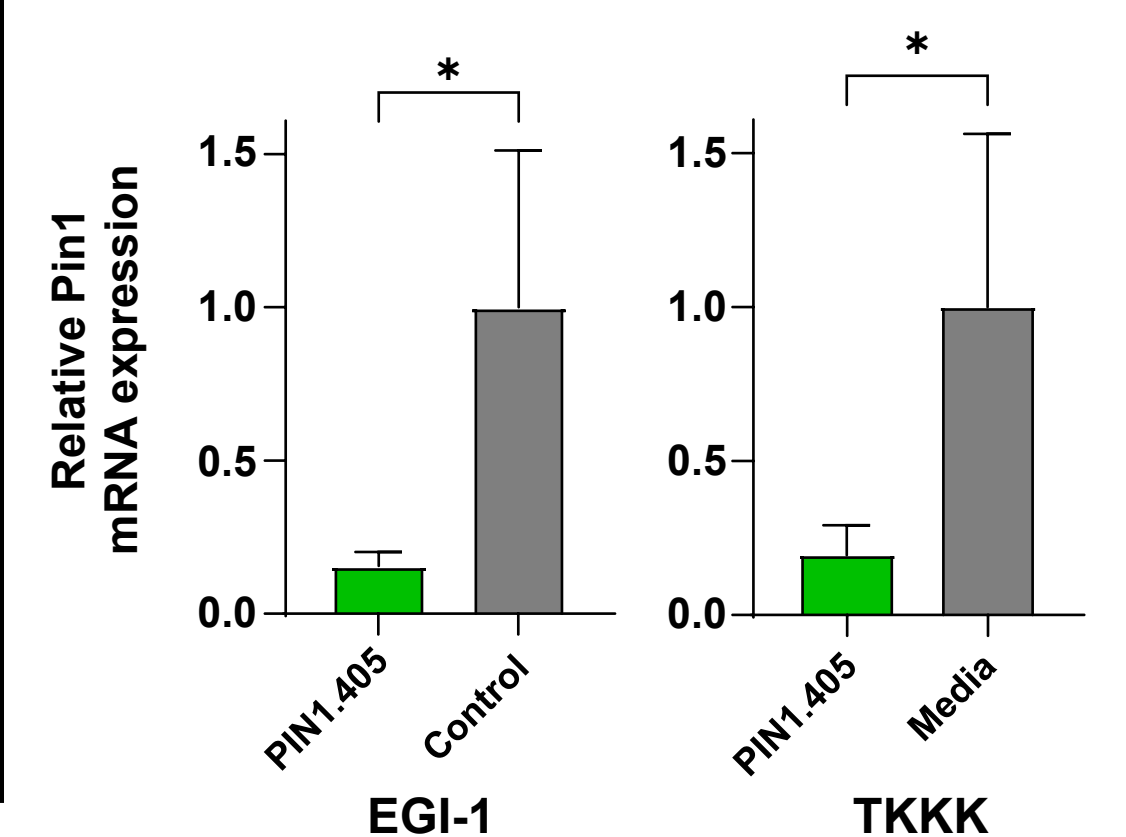
## Pin1 siRNA Screening

### Design of siRNA-L<sub>2</sub>

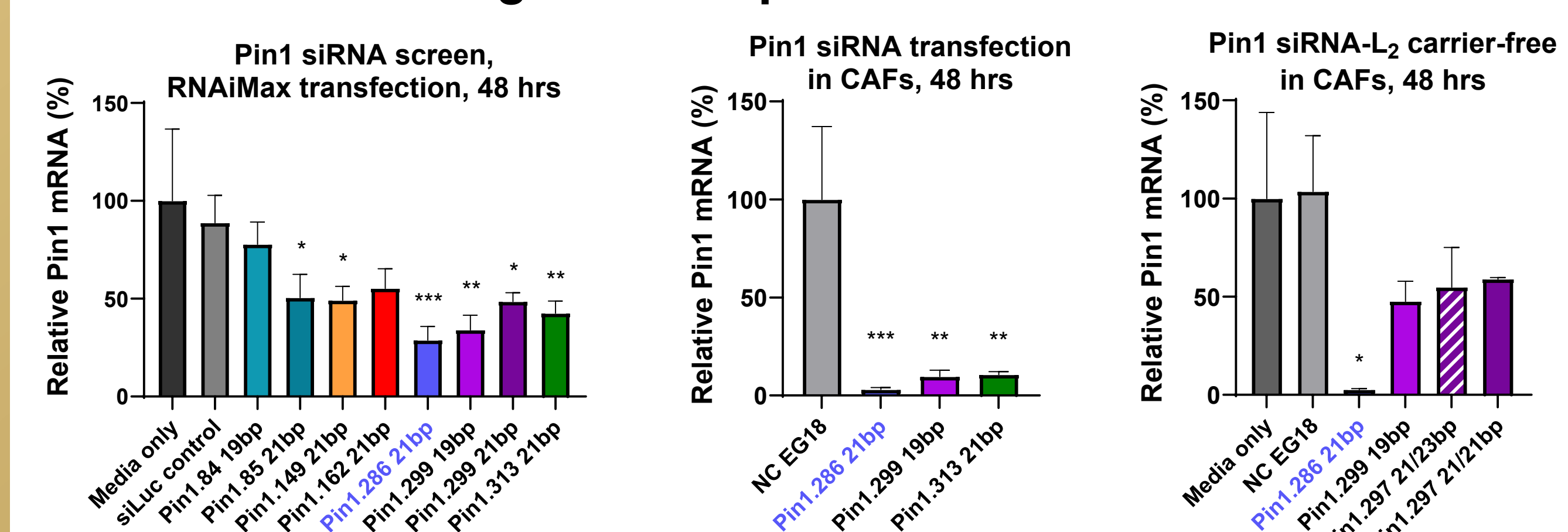
- siRNA-L<sub>2</sub> conjugates have twin lipid tails that non-covalently bind albumin, improving circulation time and tumor-tropic delivery<sup>6</sup>
- Reversible albumin binding prevents scavenger-mediated recycling
- Entirely solid-phase synthesis



### Human PIN1 silencing in CCA lines



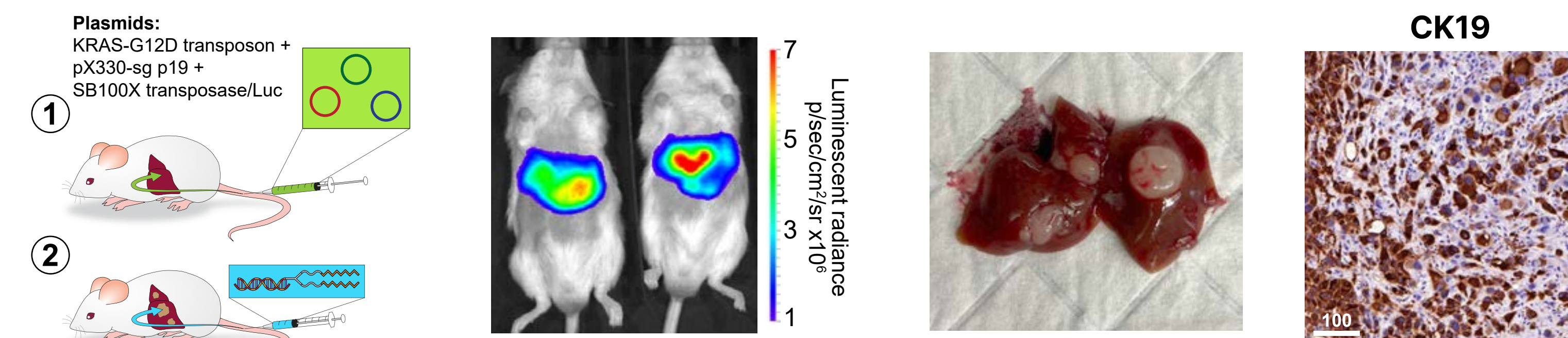
### Screening siRNA sequences for mouse Pin1



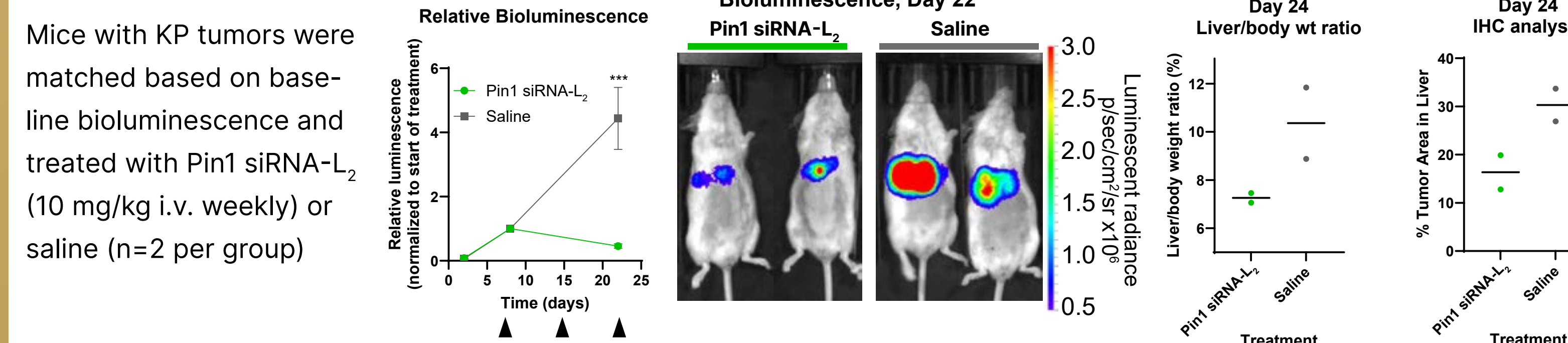
For knockdown of murine Pin1, the Pin1.286 21bp sequence results in the best knockdown, both as transfected siRNA and as carrier-free siRNA-L<sub>2</sub>.

## Pin1 siRNA-L<sub>2</sub> in Transgenic Models of CCA

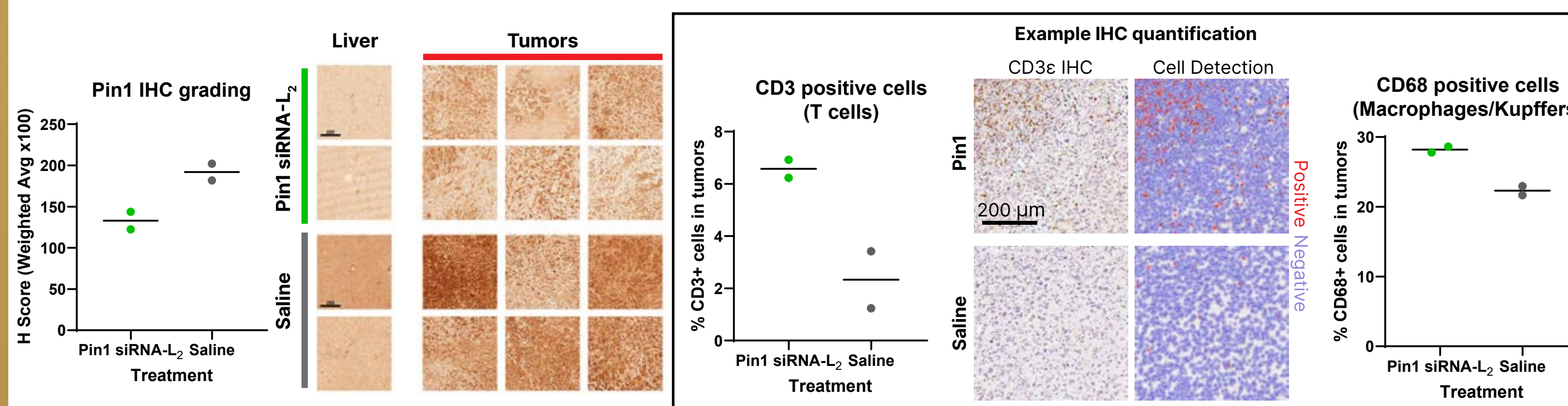
### KRAS-p19 (KP) immunocompetent mouse model of intrahepatic CCA<sup>7</sup>



### Pilot study of Pin1 siRNA-L<sub>2</sub> monotherapy in vivo



Tumor burden is reduced, as measured by *in vivo* imaging (IVIS), liver/body wt ratio, and quantification of tumor area on cross-sectional H&E



Preliminarily, Pin1 siRNA-L<sub>2</sub> treatment leads to reductions in Pin1 expression by IHC and increased percentage of T cells and, to a lesser degree, macrophages/Kupffer cells within the tumor

## Conclusions & Future Directions

### Conclusions:

- We have identified potent siRNA sequences for knockdown of human and murine Pin1
- siRNA-L<sub>2</sub> distributes into CCA tumors, including tumor and stromal cells
- Pin1 siRNA-L<sub>2</sub> is well-tolerated at therapeutic dosing in healthy mice
- Preliminarily, Pin1 siRNA-L<sub>2</sub> reduces tumor burden in a transgenic model of CCA, with trends towards reduced Pin1 expression and increased infiltration of T cells
- Pin1 staining of annotated human CCA tumor microarray shows that a subset of CCA tumors overexpress Pin1, but further analysis is needed to better understand the prognostic implications of Pin1 overexpression

### Ongoing and Future Work:

- Study underway comparing control, gemcitabine/cisplatin/anti-PD-L1 chemoimmunotherapy,<sup>8</sup> Pin1 siRNA-L<sub>2</sub>, and Pin1 siRNA-L<sub>2</sub>+chemoimmunotherapy in male and female KP mice.
- Single cell RNASeq to determine levels of mRNA knockdown in specific cell types
- CD8 depletion studies to determine role of cytotoxic T cells in mediating effects of Pin1 silencing

## References and Acknowledgments

### References:

- Lu Z and Hunter T. *Cell Research*, 2014
- Koikawa K et al. *Cell*, 2021
- Liu J et al. *Nature Communications*, 2022
- Sirica AE and Gores GJ. *Hepatology*, 2015
- Lu KP and Zhou XZ. *Nature Reviews Molecular Cell Biology*, 2007
- Hoogenboezem EN et al. *Nature Communications*, 2024.
- Zhang M et al. *Cancer Discovery*, 2023.
- Oh DY et al. *NEJM Evidence*, 2022.

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