



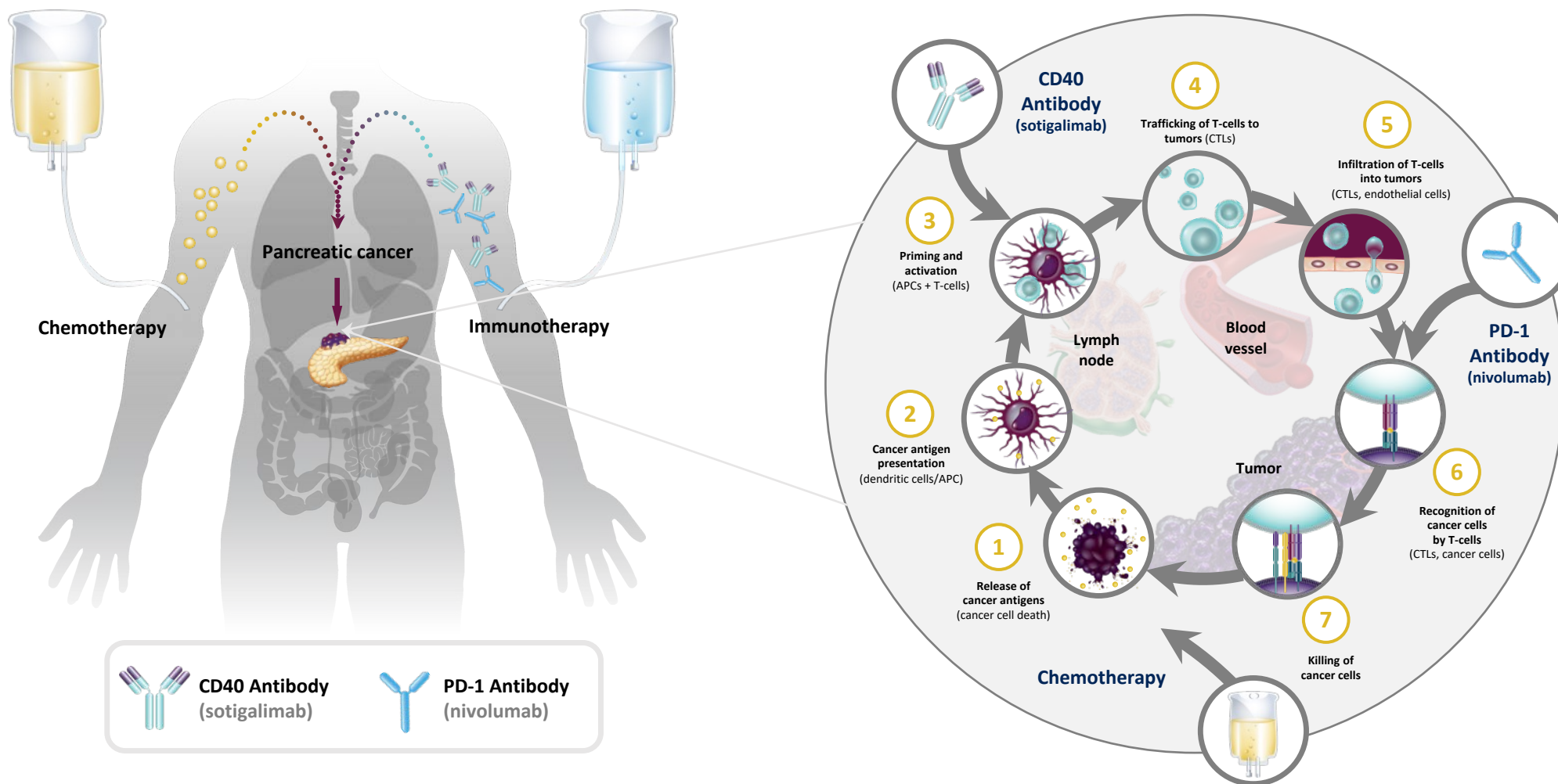
# Distinct biosignatures associate with survival after chemoimmunotherapy in a randomized, three-arm phase II study in patients with metastatic pancreatic cancer

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# PRINCE clinical trial of chemoimmunotherapy combinations in mPDAC

Gemcitabine + nab-Paclitaxel +/- CD40 agonist (sotigalimab) +/- PD-1 blockade (nivolumab)



# PRINCE trial design

1L Metastatic  
PDAC  
N=105

Ph2 Randomization

nivo/chemo (n=34): gem/nab-P\* + nivolumab 240 mg q2w

sotiga/chemo (n=30 + n=6 ph1b): gem/nab-P\* + sotigalimab 0.3 mg/kg q4w (D3 of each cycle)

sotiga/nivo/chemo (n=29 + n=6 ph1b): gem/nab-P\* + sotigalimab 0.3 mg/kg q4w (D3) + nivolumab 240 mg q2w

Follow up for  
Progression and  
Survival

\* gem/nab-P dosing and administration per SOC

Baseline and on-treatment tumor biopsies, stool collection and peripheral blood collection (including at PD)

## Key Eligibility Criteria

- 1L metastatic (recurrent or de novo) pancreatic ductal adenocarcinoma (mPDAC)
- ECOG 0-1
- Mandatory baseline tumor tissue collection
- No active CNS metastases

## Primary Endpoint

- 1-year OS compared to historical control (35%, Von Hoff 2013)

## Key Secondary Endpoints

- Objective Response Rate (RECIST 1.1)
- Duration of Response
- Progression Free Survival
- Safety

## Biomarker analyses

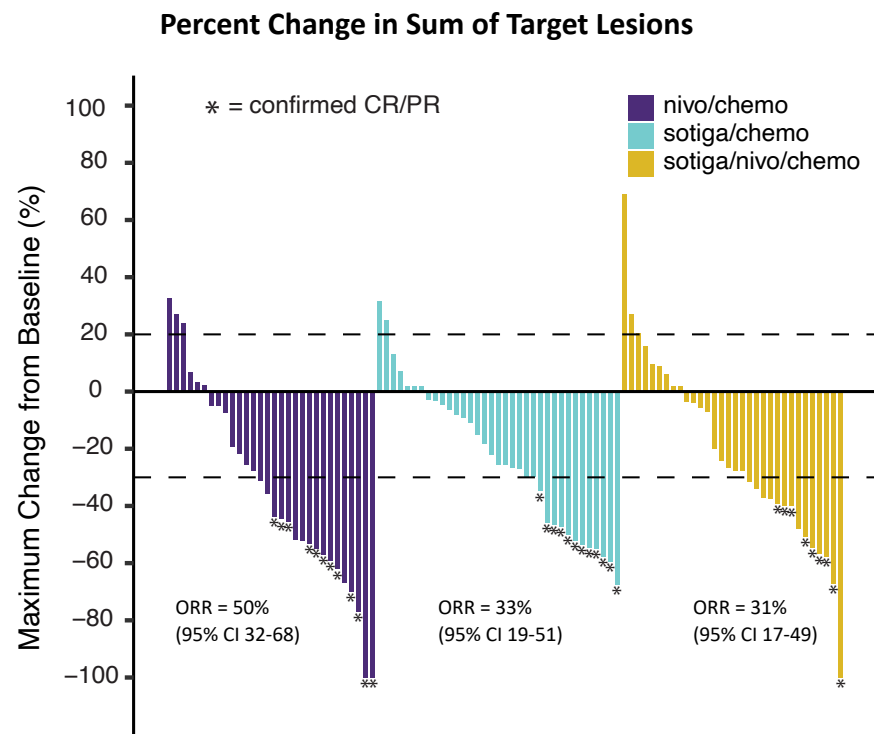
- Multi-omic, multiparameter analysis of tissue, blood and stool

## Participating Institutions

- PENN
- MSK
- MDACC
- UCSF
- UCLA
- Stanford
- DFCI

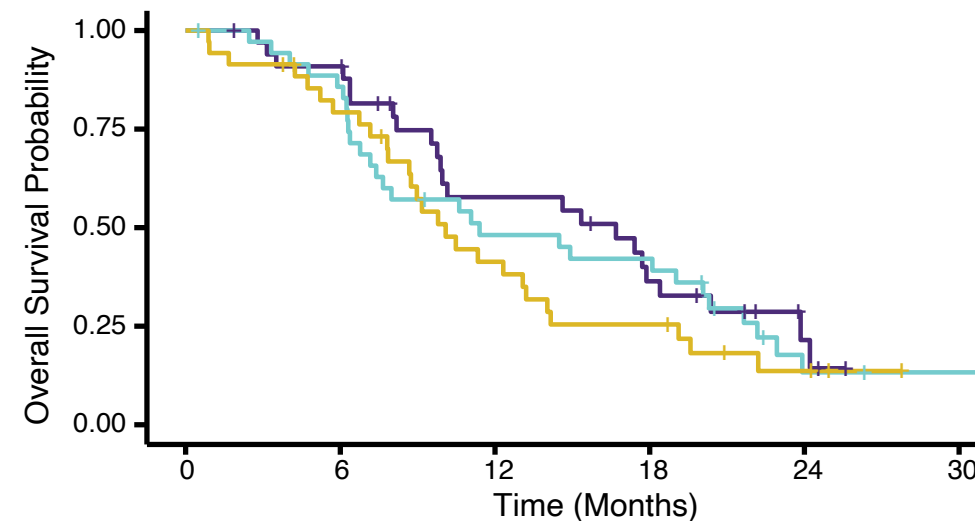
Phase 1b: O'Hara et. al, *Lancet Oncology*, 2021

# Clinical results



- No updates to clinical data presented at ASCO 2021
- Baseline characteristics were generally balanced across arms
- Treatment regimens were tolerable with manageable safety profiles
- 1-Year OS rates were compared to 35% historical control (Von Hoff, 2013)

## Overall Survival

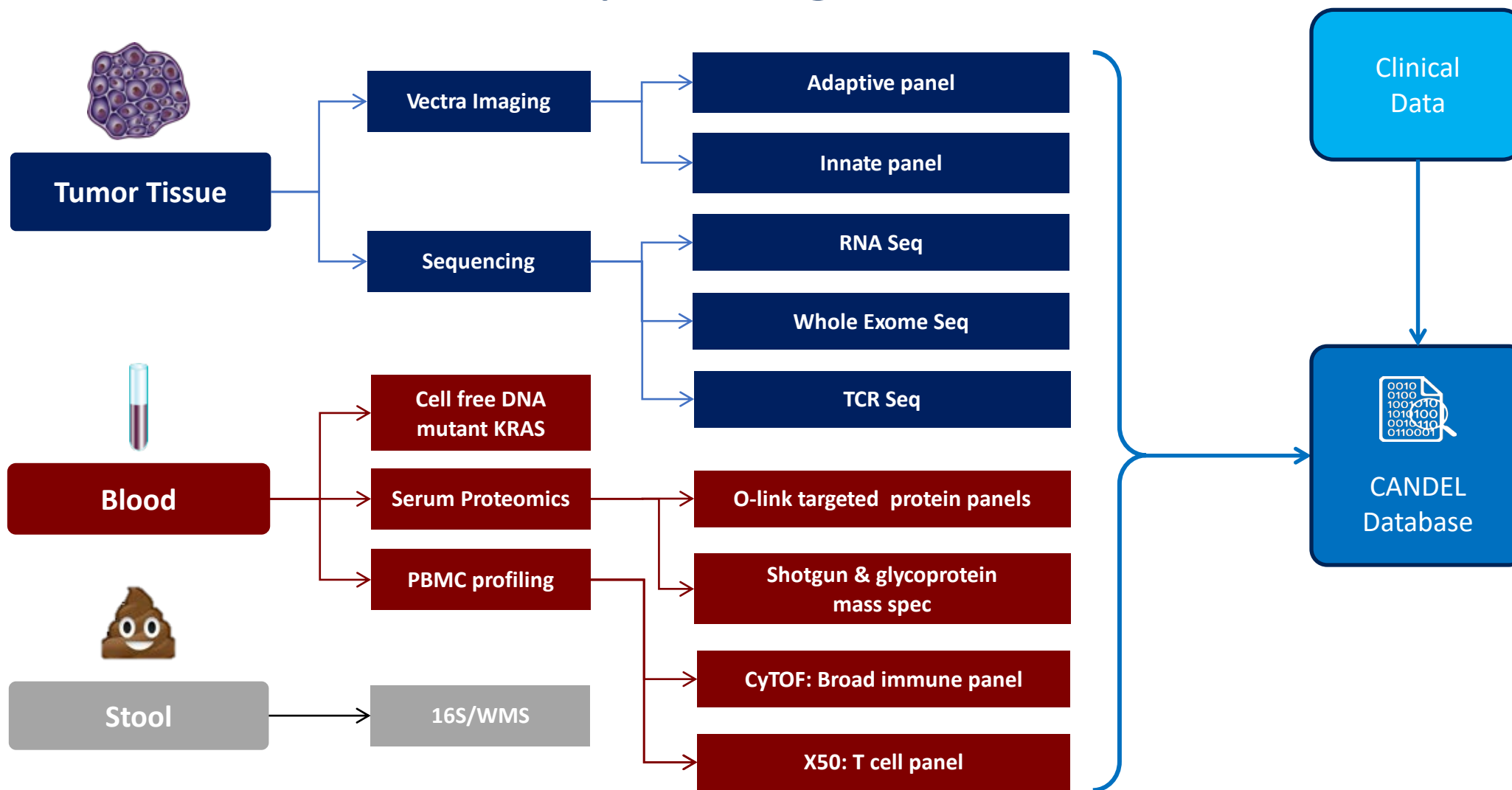


Number at risk (number censored)

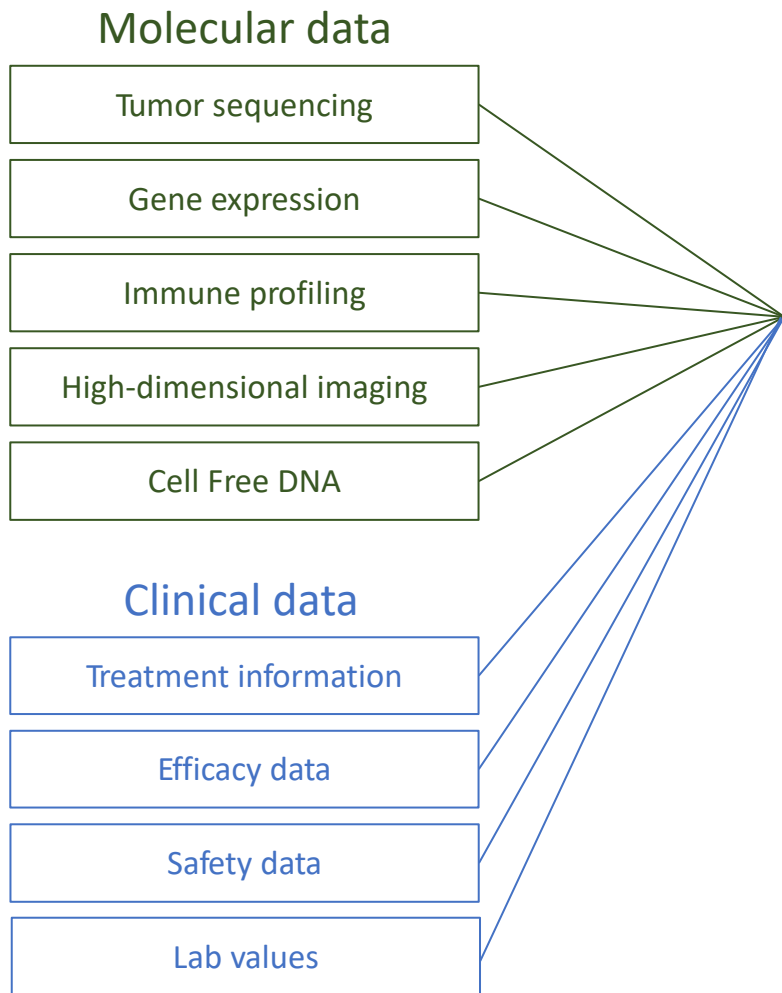
|                   | 0      | 6      | 12     | 18     | 24    | 30     |
|-------------------|--------|--------|--------|--------|-------|--------|
| nivo/chemo        | 34 (0) | 30 (1) | 17 (4) | 10 (5) | 3 (9) | 0 (11) |
| sotiga/chemo      | 36 (0) | 30 (1) | 16 (2) | 14 (2) | 3 (5) | 2 (6)  |
| sotiga/nivo/chemo | 35 (0) | 26 (2) | 13 (3) | 8 (3)  | 3 (5) | 0 (8)  |

|                   | N  | 1-yr OS rate<br>(95% lower CI) | P-value vs. 35%<br>control | Median OS<br>(95% CI)       |
|-------------------|----|--------------------------------|----------------------------|-----------------------------|
| nivo/chemo        | 34 | 57.7%<br>(41.7%)               | 0.006                      | 16.7 months<br>(9.8 – 18.4) |
| sotiga/chemo      | 36 | 48.1%<br>(33.7%)               | 0.062                      | 11.4 months<br>(7.2 – 20.1) |
| sotiga/nivo/chemo | 35 | 41.3%<br>(27.0%)               | 0.233                      | 10.1 months<br>(7.9 – 13.2) |

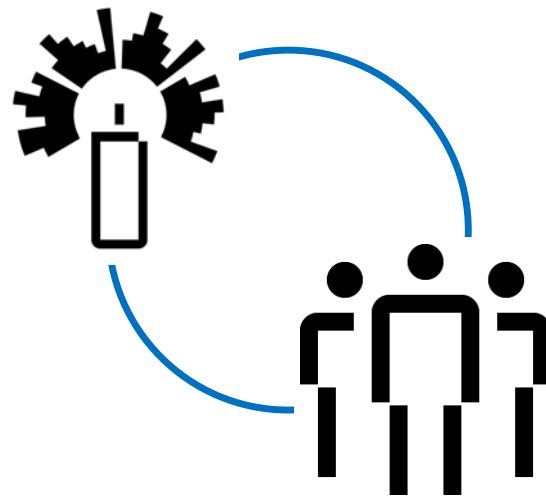
# PRINCE translational profiling



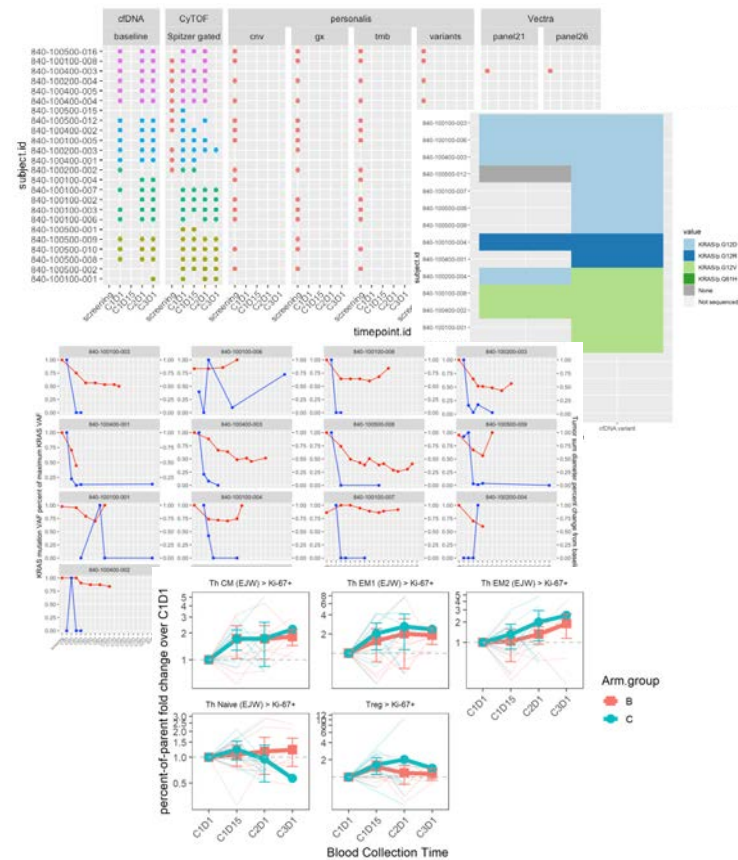
# CANDEL: Our software platform for multi-omic data analysis



**CANDEL**



**Informatics and Translational Teams**



- Biomarkers of response
- Machine learning to predict outcomes
- Insights into resistance

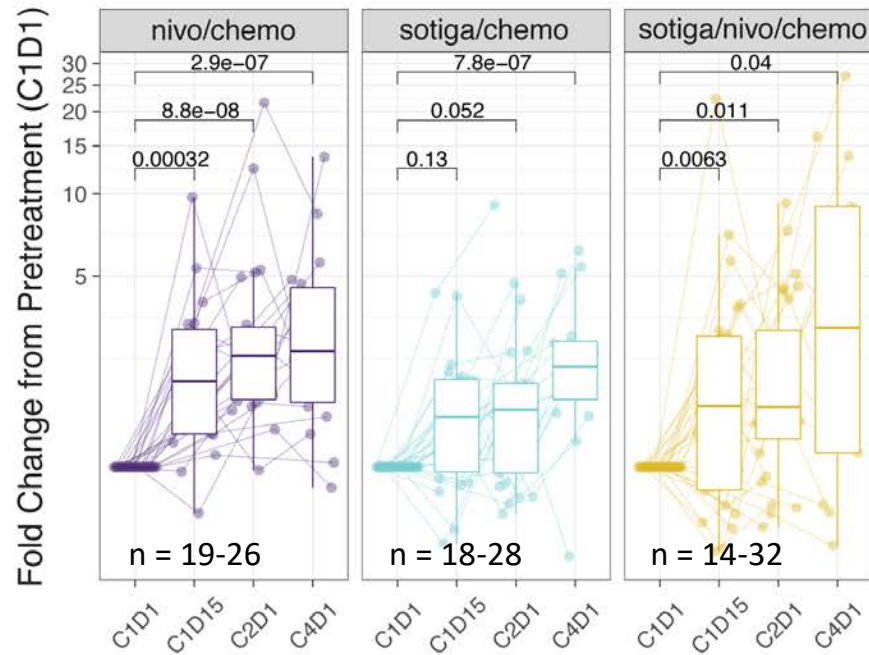
# Pharmacodynamic effects of immunotherapies



Blood

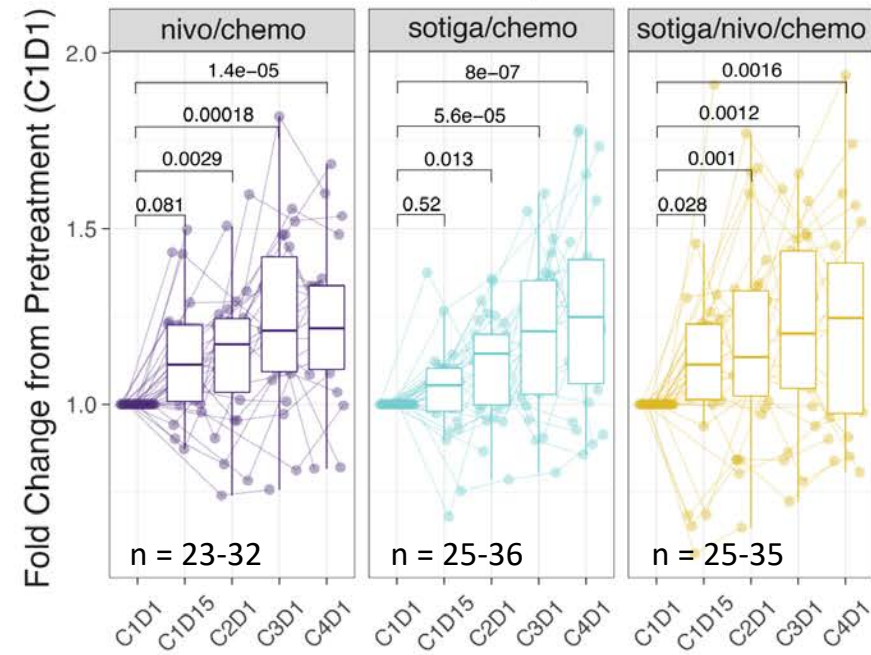
Increases in activated and proliferating T cells

% Ki-67 non-naïve CD8 T cells



Increases in soluble proteins indicating immune activation

IFN- $\gamma$

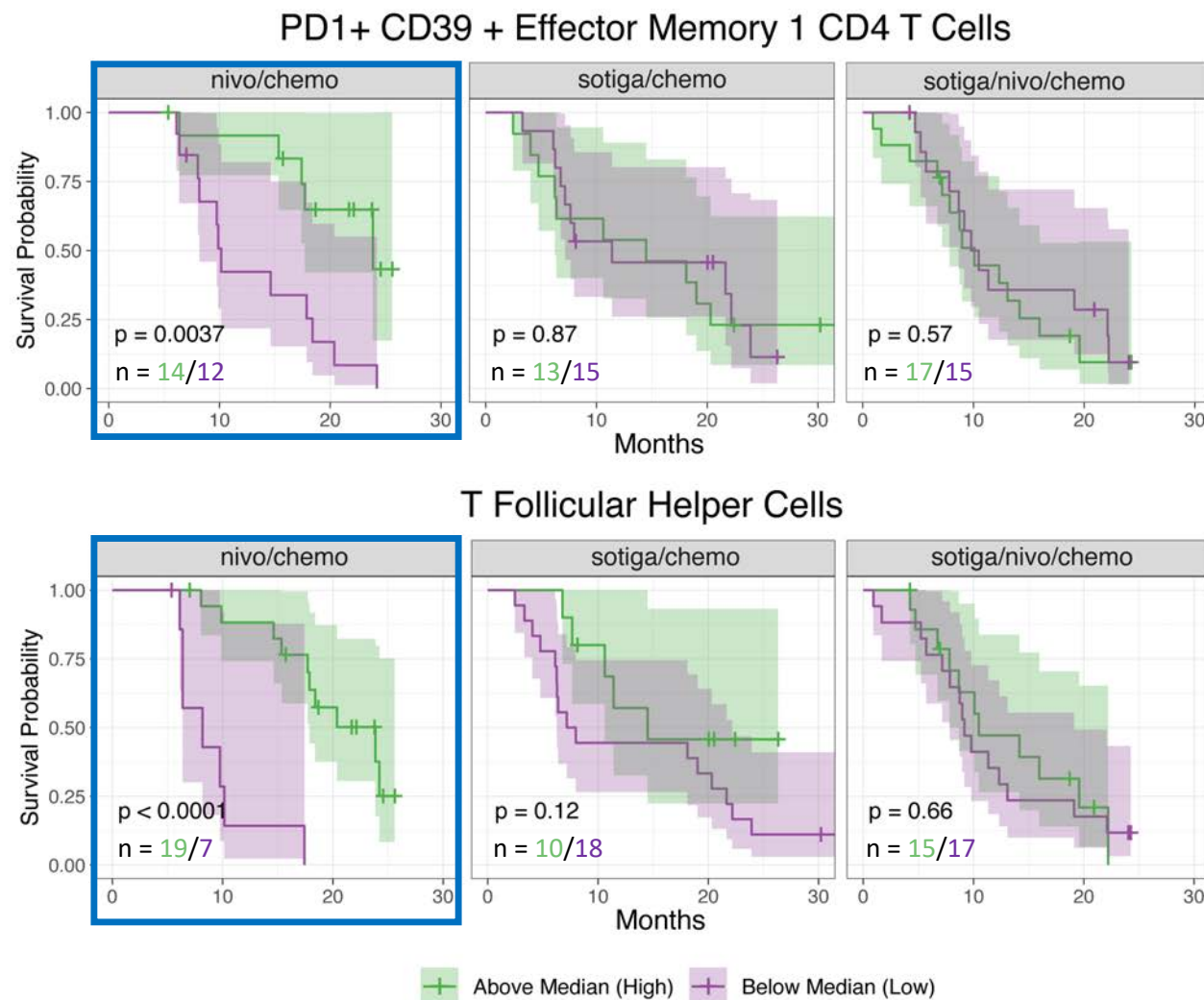


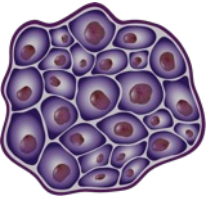
# Circulating biomarkers of survival following nivo/chemo



Blood

- Baseline levels of T cell populations in circulation predict survival following nivo/chemo
- Many predictive T cell populations show an activated, antigen-experienced phenotype
- Biomarkers predict survival after controlling for clinical covariates

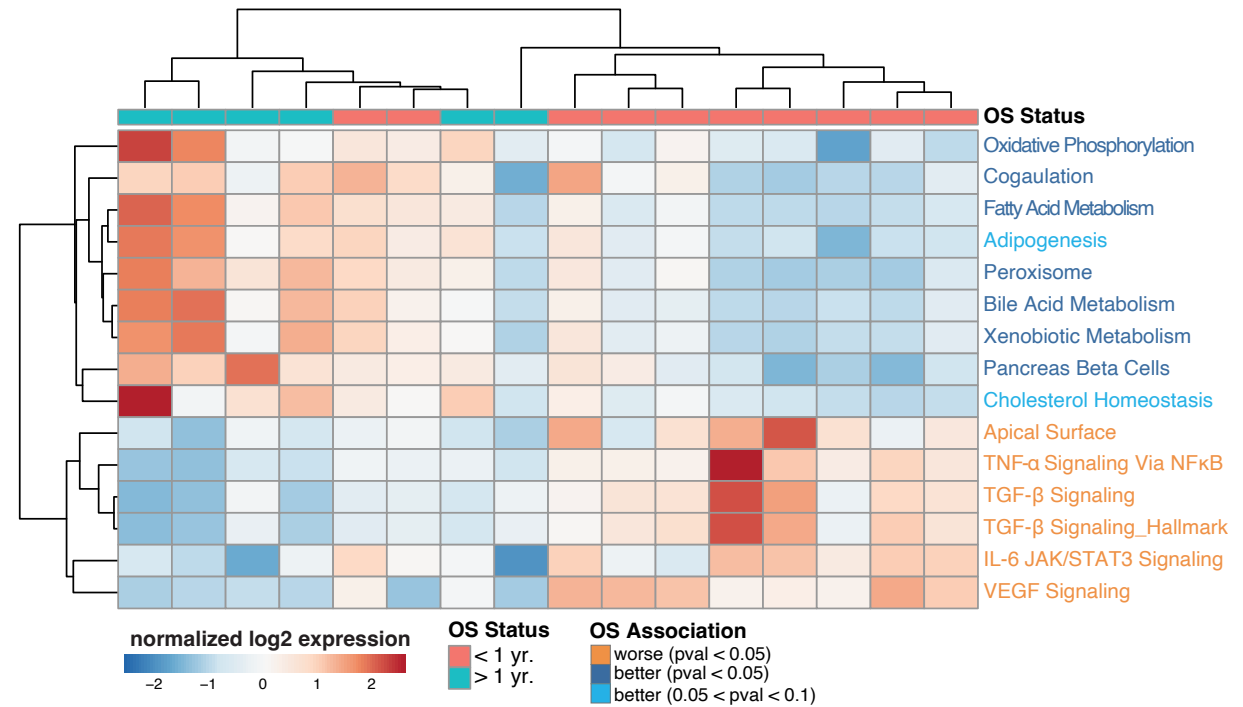




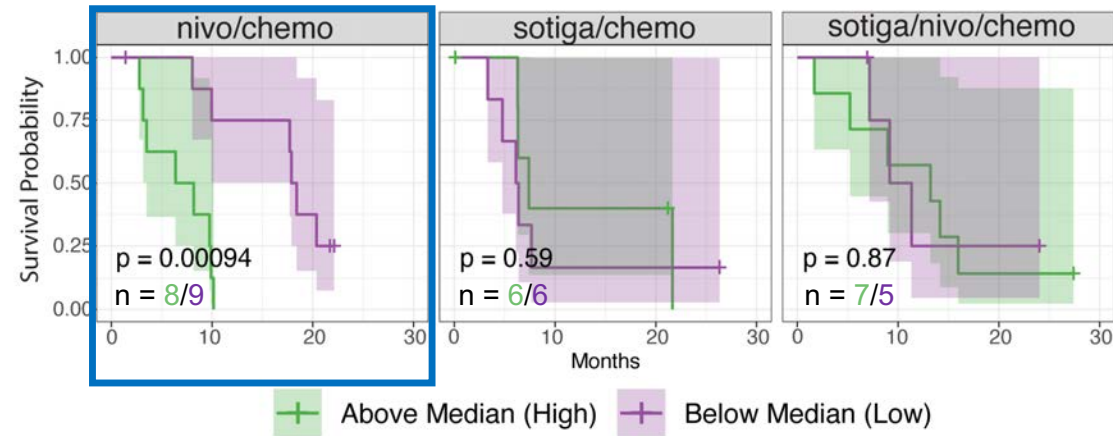
Tumor

# Tumor biomarkers of survival following nivo/chemo

- Several baseline tumoral gene expression signatures correlate with survival following nivo/chemo
- Signatures reflect biology of a tumor low in immune suppression and in a favorable oxidative lipogenic metabolic state



## TNF-α Signaling via NFκB



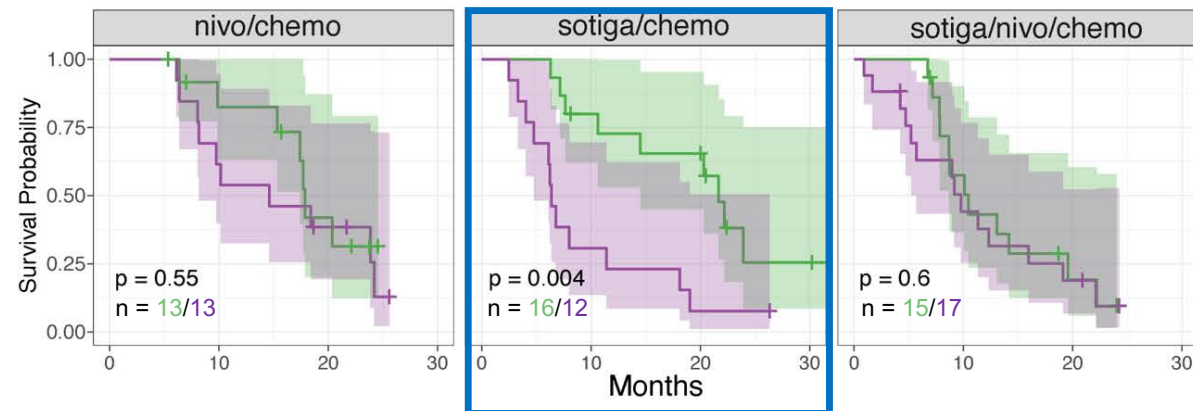
# Circulating biomarkers of survival following sotiga/chemo



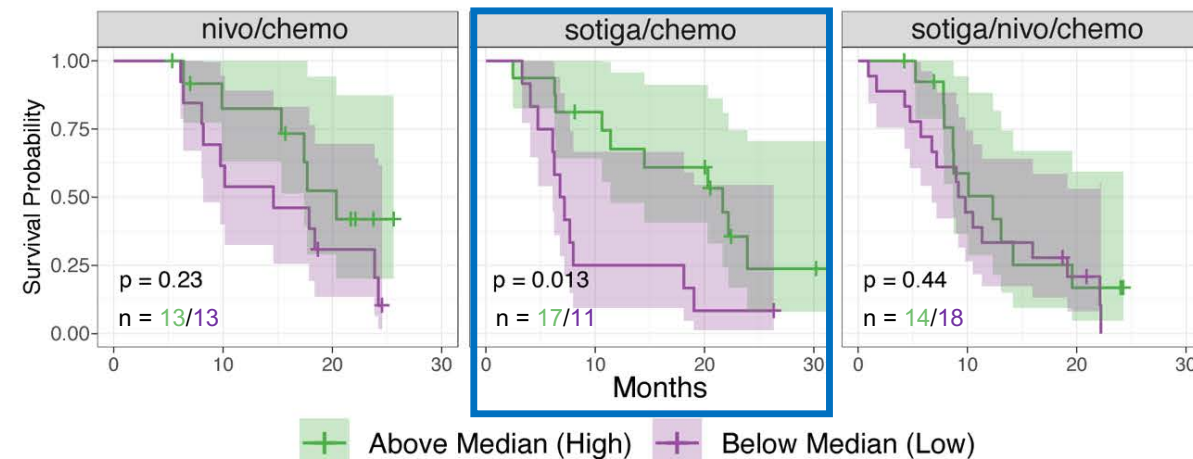
Blood

- Pretreatment levels of CD4 T cell, B cell, and dendritic cell subsets predict survival benefit
- Predictive cell populations reflect the mechanism of action of agonistic CD40 therapy
- Biomarkers predict survival after controlling for clinical covariates

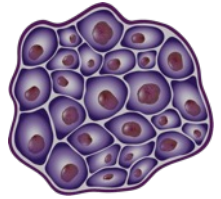
## PD-1+ Tbet+ non-naïve CD4 T cells



## Tbet+ Eomes+ non-naïve CD4 T cells

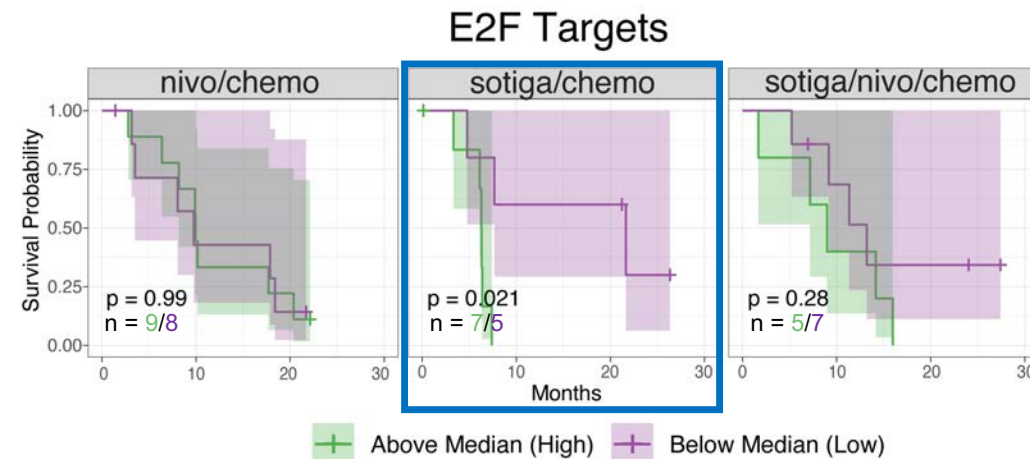
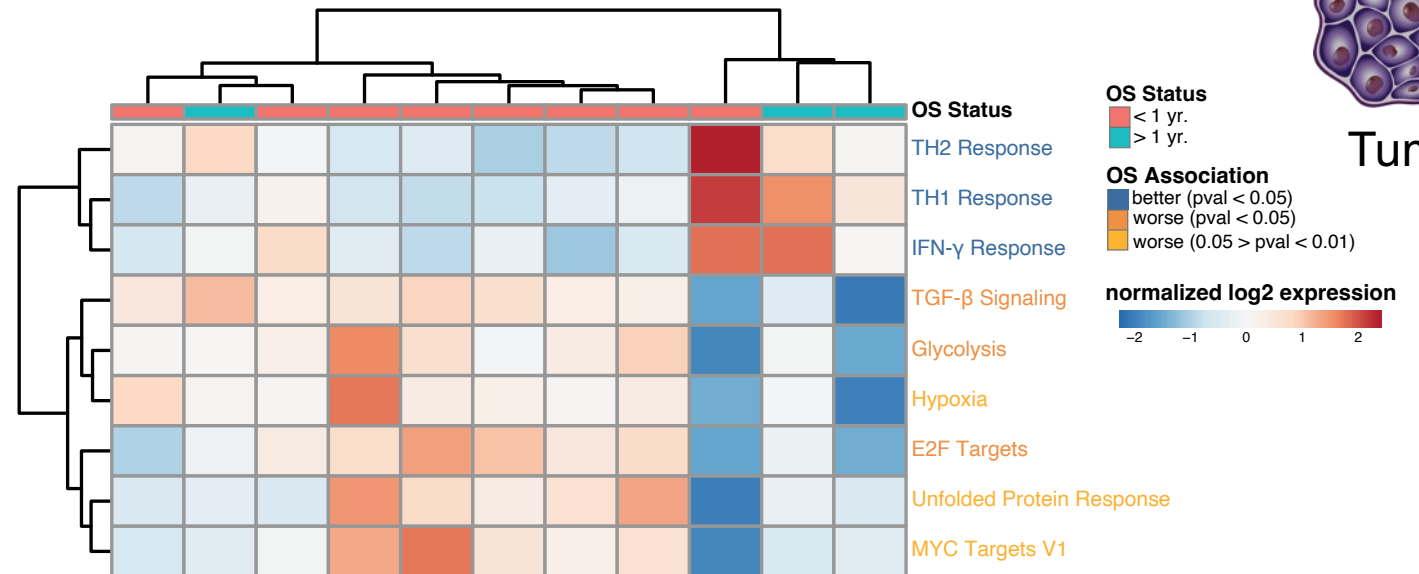


# Tumoral biomarkers of survival following sotiga/chemo



Tumor

- Pretreatment tumoral gene expression signatures and immune cell populations correlate with survival benefit
- Longer surviving patients have tumors higher in non-proliferating helper CD4 T cell infiltration and lower in E2F and Myc signaling



# Biomarkers of survival following sotiga/nivo/chemo

- Few factors in the tumor or circulation predicted survival benefit
- Relevant biomarkers from other arms were not predictive of survival benefit
- Hyperactivation may be leading to a dysfunctional immune state

# Conclusions

- **Immuno-pharmacodynamic effects were observed in each arm**
- **Novel circulating and tumoral biomarkers predict overall survival following treatment with nivo/chemo or sotiga/chemo**
- **Prospective studies are warranted to test the validity of blood-based predictive biomarkers of response to nivo/chemo or sotiga/chemo**
- **Proof of principle for engagement of the PICI network to conduct multi-cohort, multi-center bench-to-bedside translational and informatic studies**
- **Established a framework for REVOLUTION platform in 1L mPDAC (NCT04787991)**

# Acknowledgments

## Patients & their families

Sean Parker

**Lead Study PI:** Robert H. Vonderheide

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**PICI Central:** Sultan Nawabi, Justin Fairchild, Jingying Xu, Joyson Karakunnel, Lisa Butterfield, Ramy Ibrahim, Ute Dugan, Christine Spencer, Mike Travers, Robin Kageyama, Diane Da Silva

**Sites:** Rosemarie Mick, Kate Byrne, Cecile Alanio, John Wherry, Jonni Moore, Derek Jones, Erica Carpenter, Jacob Till, Susan Domchek, R. Schretzenmair, J. Jakobowski, G. Brake-Silla, Arvin Ruiz, Y. Li

**Partners:** Vanessa Hubbard-Lucey, Marco Tognetti, Lisa Salvador, Charles Abbott, Lee McDaniel, Sean Boyle, Kamil Sklodowski, R. Bruderer, L. Reiter

## Funding:



## Study drug:

