Development of a HIF2a-Targeted RNAi Therapeutic for the Treatment of ccRCC

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Outline

- Development of ARO-HIF2
- Key pre-clinical proof-of-concept results
- ARO-HIF2 clinical study
Arrowhead’s ARO-HIF2 Program

- Second Gen Platform: **Targeted RNAi Molecules (TRiM™)**
- Enhanced delivery via an integrin receptor (αvβ3) that is over-expressed in many cancers
  - Tumor tissue microarrays confirmed receptor expression in ccRCC at high frequency
- RNAi trigger specifically targets HIF2α mRNA
  - Over-expression in ccRCC especially with VHL mutations
    - HIF2α is regarded as a key tumorigenic driver of ccRCC
  - Limited restrictive expression in normal tissues
  - Chemically modified to enhance potency and prevent immune activation
  - Minimal off-target risks
Tumor Delivery is Ligand Dependent

**No ligand**

**With Ligand**

- Efficient delivery to all tumor cells
- Weak delivery without ligand

2 mg/kg Cy3-labeled ARO-HIF 4 h after injection

- **Red** = ARO-HIF2
- **Blue** = nuclei
- **Green** = actin (cell membrane)

A498 ccRCC orthotopic tumor mouse model
A498 orthotopic kidney xenograft mouse model

- A498 is an established ccRCC cancer cell line
  - VHL mutated, HIF2a over-expressed
  - Integrin αvβ3 positive

- SEAP-A498 model
  - Stably expresses SEAP (secreted embryonic alkaline phosphatase)
  - Good correlation between SEAP levels and tumor volumes

- Sensitive serum biomarker to monitor tumor growth

![Tumor volume vs. SEAP levels](image-url)
ARO-HIF2 Dose Response in A498 Mouse Model

ARO-HIF2 Dose Response (single injection)

- Single dose on study Day 1
- Gene expression (KD) on Day 8
- Shallow dose response above 6 mg/kg
ARO-HIF2 Response Duration in A498 Mouse Model

- Single dose on study Day 1
- Nadir Day 8, HIF2α 82.2 % KD
- Max KD last for about 1 week
Reduction in HIF2α mRNA and Protein

Single 10 mg/kg iv dose (n= 4 to 5)
Tumor HIF2α mRNA and protein levels one week after dosing
Multi-Dose Tumor Growth Inhibition (TGI) Study

- Eight weekly doses of 13 mg/kg or 26 mg/kg of ARO-HIF2
- Weekly SEAP monitoring for TGI
- End of study tumor HIF2α gene silencing, sizes and histology
ARO-HIF2 TGI Study: Response by SEAP

- A498 orthotopic SEAP mouse model
- Similar TGI response based on SEAP readout
- Both treatment groups had mice showed regression by SEAP
- One mouse in 26 mg/kg treatment group showed sign of treatment escape by SEAP readout
A498 TGI Study: Response by Tumor Volume and Gene Silencing

- Treatment groups show better BW maintenance.
- Both dose levels showed strong tumor growth inhibition (TGI) and deep HIF2α mRNA KD.

**Body Weight (g)**

- Saline
- 1x 13 mg/kg, weekly
- 1x 26 mg/kg, weekly

**Tumor Volume**

- Symbols with lighter tone: escapee by SEAP
- All graphs show mean (SEM)

**HIF2α mRNA KD**

- 86.2%
- 91.1%

D37 euthanized 1 mouse
Tumor Histology

Day 1 baseline
Day 57 Saline
Day 57 13 mg/kg
Day 57 26 mg/kg

- ARO-HIF2 treated group showed wide-spread tumor destruction
- Loss of clear cell characteristic
- Areas of apoptosis and necrosis

→ Necrosis
▼ Macrophage infiltration
⇒ Apoptosis
Summary of ARO-HIF2 Pre-Clinical Studies

• Efficient ligand enhanced tumor delivery of ARO-HIF2
  • Demonstrated deep HIF2α mRNA knockdown with corresponding reduction in HIF2α protein in tumor cells
  • Inhibition of tumor growth and improved overall survival in tumor models

• Single and multiple doses up to 26 mg/kg tested

• Exploratory toxicity studies in rats and NHPs up to 120 mg/kg
  • No drug-related observations or effect on body weight
  • Minimal LFT increase at top dose
  • No decrease in EPO level
  • No significant HIF2α mRNA KD in NHP liver, kidney, spleen and adrenal at 80 mg/kg
ARO-HIF2 Phase 1 Clinical Program

- Phase 1 dose range finding study
  - To be conducted in I/O and/or anti-VEGF refractory ccRCC patients
- Initiated 08/2020, dose escalation on-going
- Primary objectives:
  - Safety & determination of phase 2 dose
- Secondary objectives
  - PK, efficacy based on RECIST
- Key exploratory objective
  - Tumor biopsy HIF2α expression
Thank You