



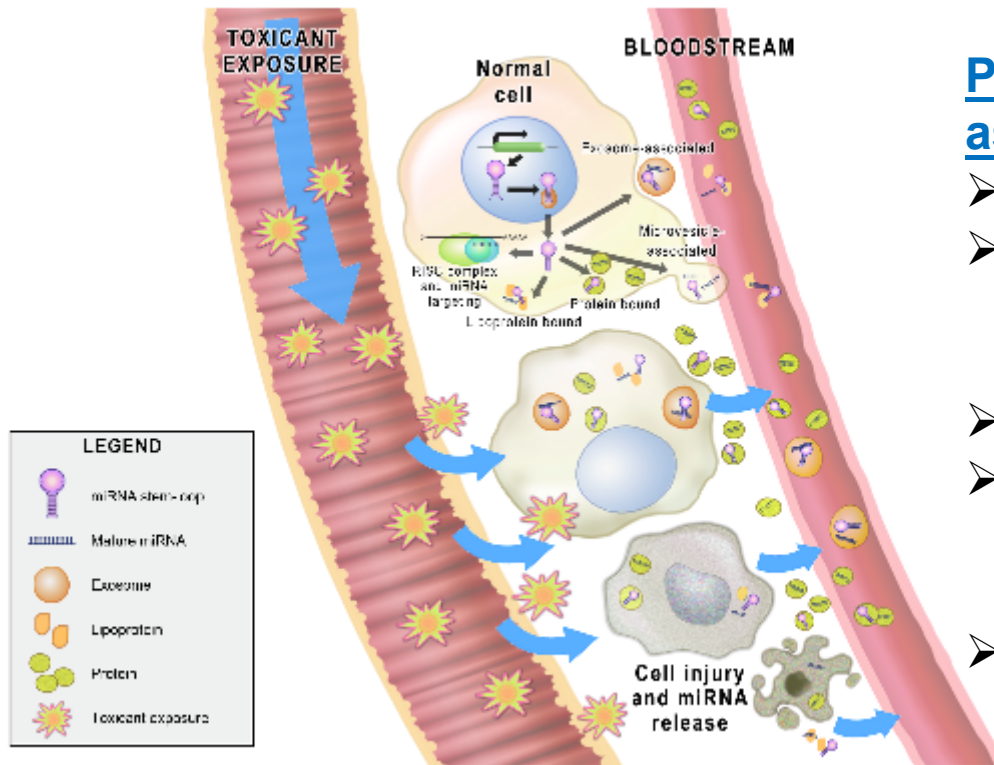
# **The rat microRNA body atlas: miRNA identification and characterization**

**A Project of the HESI Genomics Committee**

**Alison Harrill, Aaron Smith, and Raegan O'Lone**

# MicroRNAs: Tissue Specific Biomarkers

miRNA function: RNA silencing and post-transcriptional regulation



## Potential advantages of miRNAs as biomarkers:

- **Sensitivity and specificity.**
- **Can easily measure**, utilizing a small volume of serum in a qPCR based format.
- **Stable** in serum and plasma.
- **Conservation across species** and non-Ab detection enables ready translational use.
- Identification of **isomiRs** may yield **additional candidate biomarkers.**



Fill gaps for tissues  
where no reliable injury  
biomarkers exist



Decrease time and cost of  
drug development



## **Opportunities for miRNA Biomarkers in Chemical/Drug Safety Assessments**



Enhance clinical translation  
& clinical monitoring



Improved sensitivity and  
specificity over existing  
biomarkers





# Rat miRNA Body Atlas Phase I – Identification of Tissue Specific miRNA

## 23 Tissues Assessed

Kidney, whole	Duodenum
Medulla	Cerebrum
Cortex	Cerebellum
Liver	Hippocampus
Heart	Brain stem
Pancreas	Dorsal root ganglion
Adrenal	Soleus
Glandular stomach	Biceps femoris
Non glandular stomach	Whole blood
Jejunum	Testis
Ileum	Ovary
	Uterus



Tissues were extracted from 5 male and 5 female SD rats and RNAs were subjected to Illumina deep sequencing.

RNA quality was assessed and found to be of good quality (pancreas was an exception with a lower than average quality not unusual for that tissue)

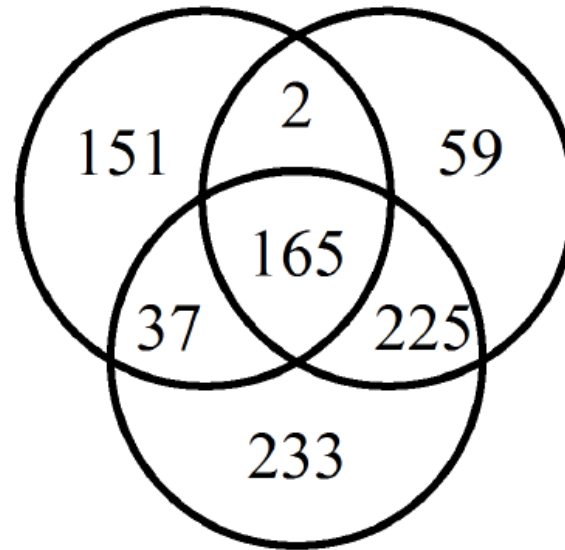


# Various Analysis Methods Were Utilized to Determine Tissue miRNA Abundance

Maastricht University

Eli Lilly

Each institution used a different statistical method for the RNA-Seq data



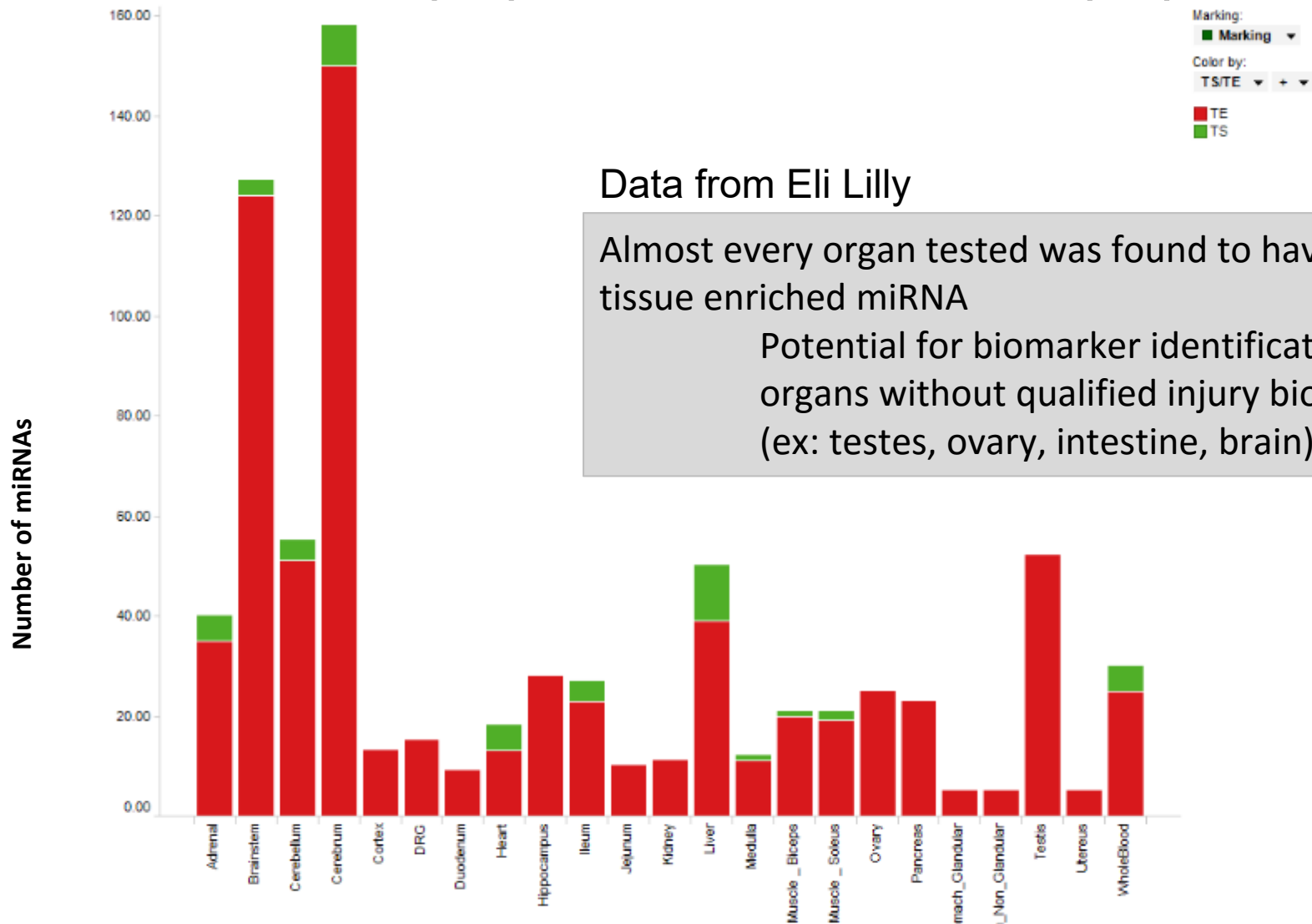
NIEHS

## Conclusions:

- Analysis method is important, 165 miRNA in agreement across methods
- A single analysis method would not have found as many miRNA with high confidence as potential biomarker candidates



# Tissue Specific (TS) and Tissue Enriched (TE) miRNAs



Data from Eli Lilly

Almost every organ tested was found to have some tissue enriched miRNA

Potential for biomarker identification for organs without qualified injury biomarkers (ex: testes, ovary, intestine, brain)

- Rat body atlas paper submitted
- Rat miRNA body atlas to be made available to other HESI committees for use in future studies



# WHAT LIES AHEAD?

- Pool of Tissue Specific & Tissue Enriched miRNAs identified
  - Methods to further characterize putative biomarker candidates?
  - The committee is scoping a 2<sup>nd</sup> phase of work...

## Rat miRNA Body Atlas Phase II

---

### 1<sup>st</sup> Round Scoping Assessment

**Characterizing 'information content' of miRNA profiles and integrating other endpoints, longitudinal analyses, adding mechanistic value, and tying to the underlying biology**



---

### 2<sup>nd</sup> Round Scoping Assessment – Under Consideration

Investigating cell or tissue region specific localization of miRNAs  
-Which tissue or cell type?

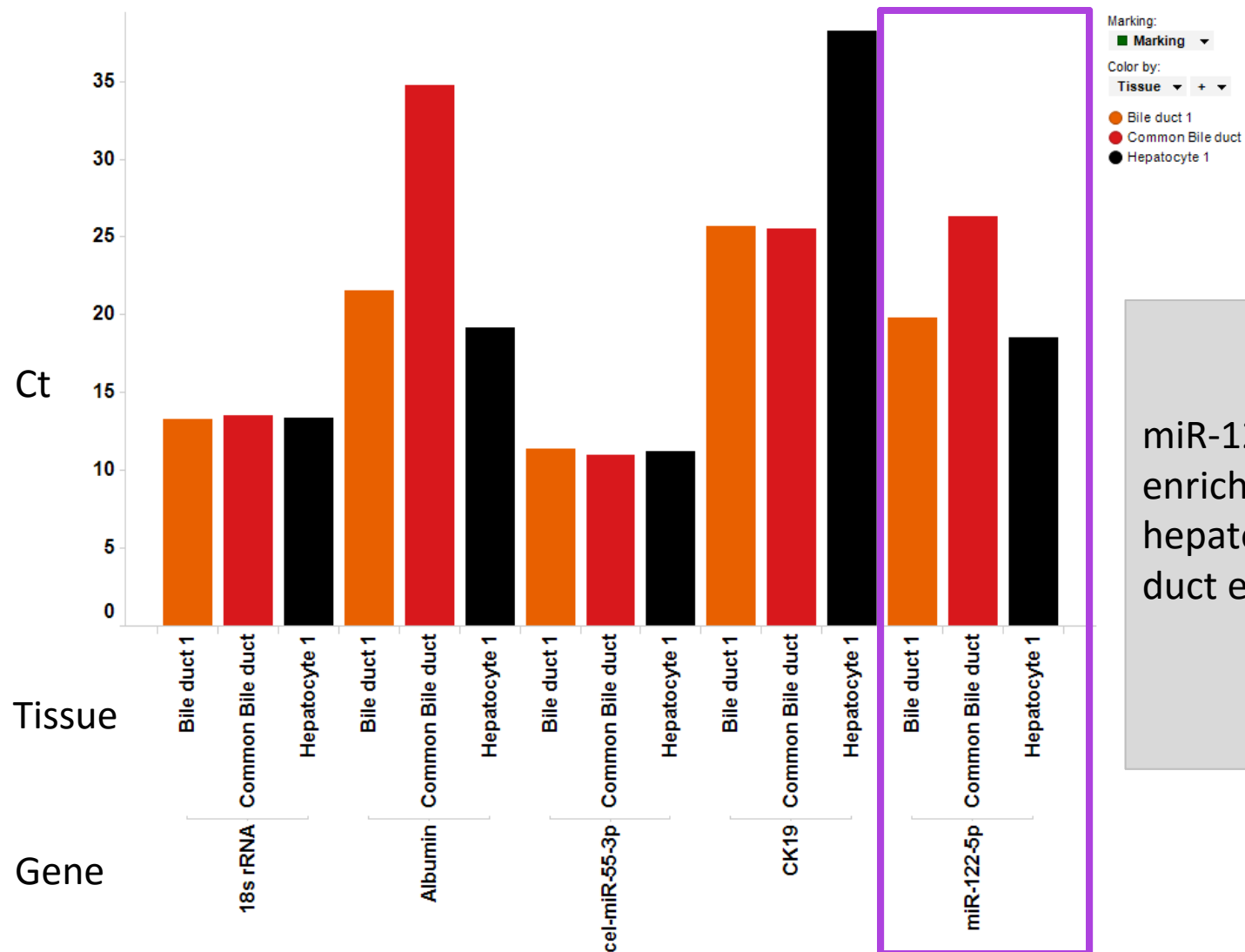
Exploring kinetics of miRNA level changes in tissue and circulation (potentially including transient vs. long-term miRNA changes)

-When is the optimal time to measure a miRNA biomarker?

---



# Cell Type Specificity: Example Data with miRNA-122



miR-122-5p is enriched ~64 fold in hepatocytes vs. bile duct epithelial cells

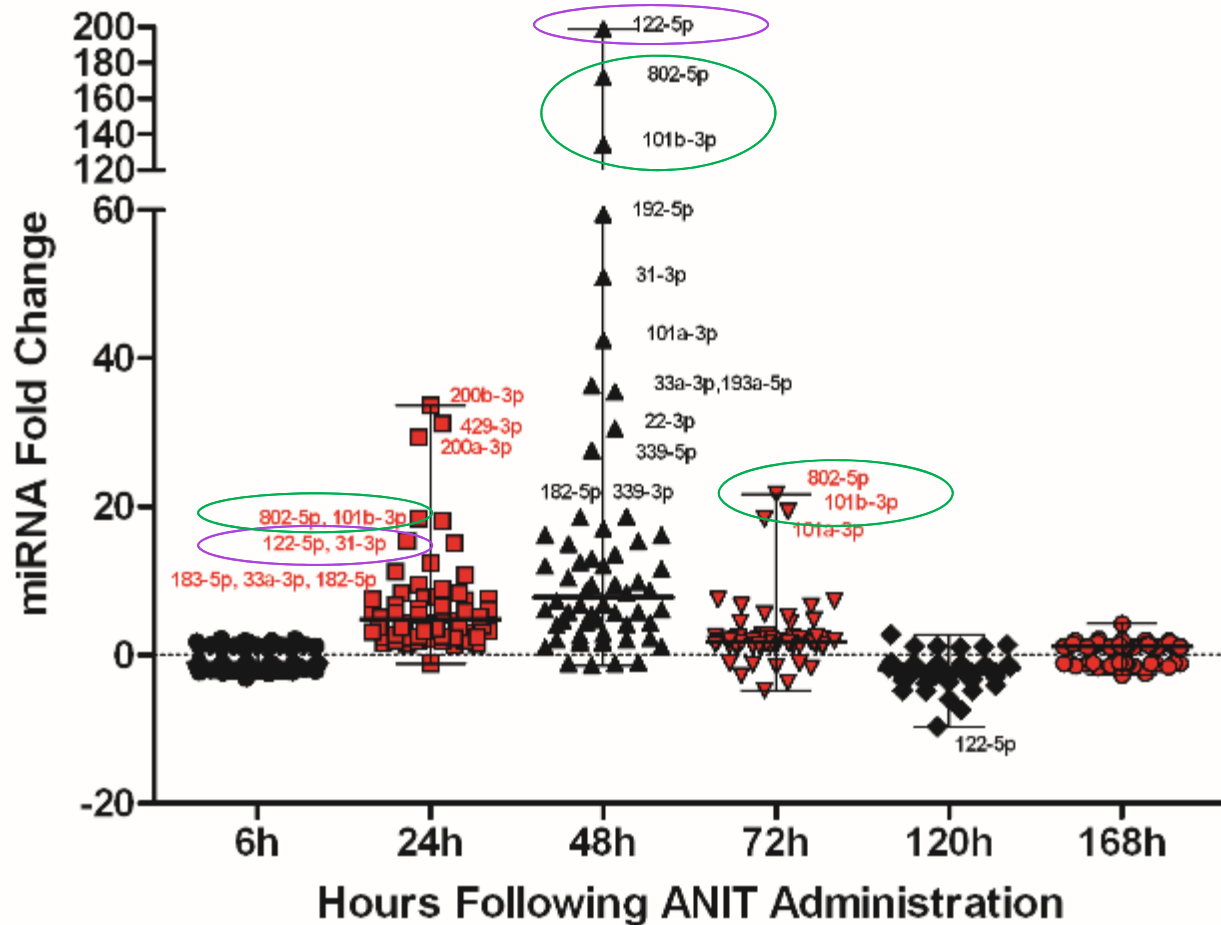
Albumin= Hepatocyte marker

CK19= Bile duct marker

Data generated by Aaron Smith, Eli Lilly



# miRNA Time Course Profiles: Plasma Following Liver Toxicity



miR-122-5p is released only during peak liver injury

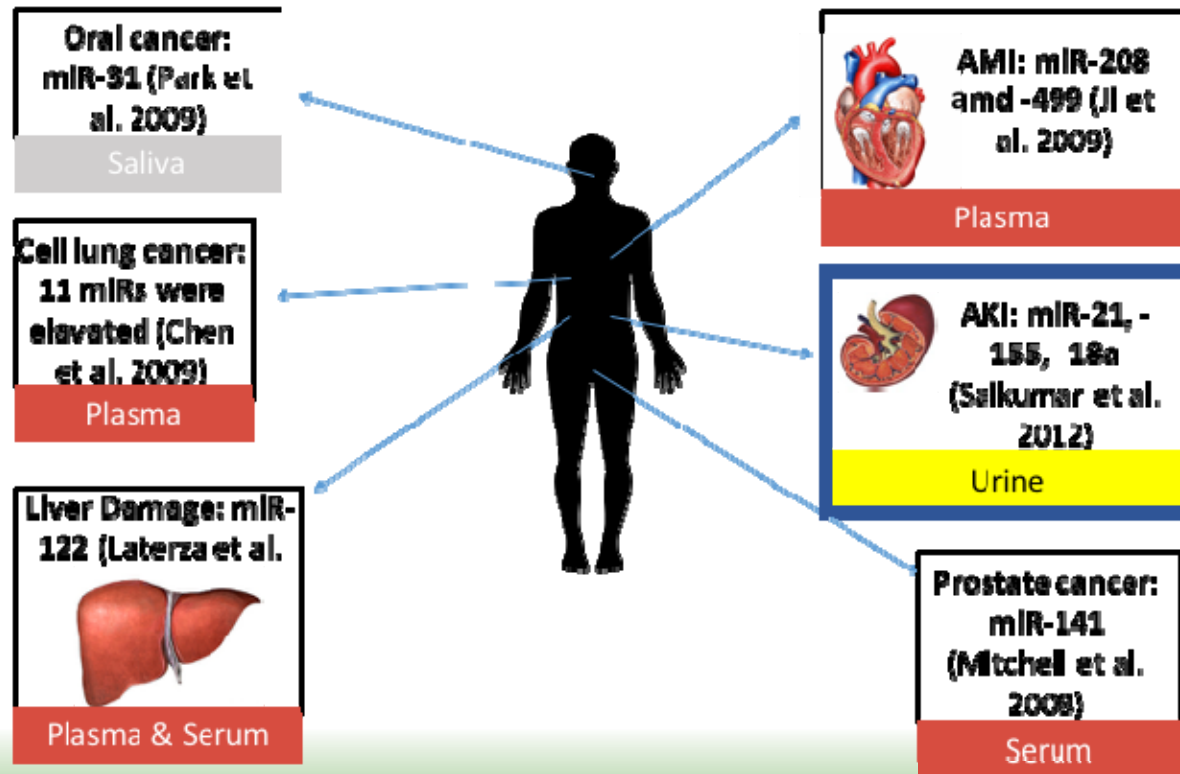
miR-802-5p and miR-101b-3p offer advantage of increases during peak injury AND repair phases

**Timing of measurement is important**



# Impact of miRNA Body Atlas

- Development of “liquid biopsies” – inform underlying pathology
- Translation of miRNA biomarkers across species
- Clinical translation for organ injuries under a variety of disease states and contexts (not exclusive to toxicology)
- Foundational research for others to interrogate their own tissue-specific questions
  - Collaboration with other HESI committees, consortia, academic institutions



# Acknowledgements

HESI Genomics Committee

**Working Group Leaders:** Aaron Smith, Karol Thompson, Heidrun Ellinger

Data Generation & Analysis:

**Eli Lilly:** Aaron Smith, John Calley, Sachin Mathur, Hui-Rong Qian, Han Wu, Mark Farmen, David Hall, David Watson

**Maastricht University:** Florian Caiment

**NIEHS:** Pierre Bushel, Jianying Li

**Takeda:** Craig Fisher, Patrick Kirby, Erik Koenig



# BACKUP SLIDES



# miR-122-5p localization in hepatocytes vs bile ducts

1. Rat liver was collagenase digested and the common bile duct (CBD) was cannulated.
2. Blue agar was injected into the CBD and was visibly detectable in the bile ducts within the liver.
3. The CBD was placed in Trizol and the hepatocytes were separated from micro-dissected liver.
4. RNA was isolated from CBD, hepatocytes and micro-dissected bile ducts.

