

# Extracellular RNA Communication

## NIH Common Fund Program





National Institutes of Health Office of Strategic Coordination - The Common Fund

### **Extracellular RNA Communication Program**

New Paradigm: RNAs are released from cells and (may) go on to influence cells that receive them. RNAs are found in the environment: food, bacteria (think microbiome).

nature

cell biology

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Exosome-mediated transfer of mRNAs and microRNAs is a novel mechanism of genetic exchange between cells

Hadi Valadi<sup>1,3</sup>, Karin Ekström<sup>1,3</sup>, Apostolos Bossios<sup>1</sup>, Margareta Sjöstrand<sup>1</sup>, James J. Lee<sup>2</sup> and Jan O. Lötvall<sup>1,4</sup>

An opportunity to explore new paradigms of cell-to-cell communication based on release, transport, uptake, and regulatory role of exRNAs



### **Extracellular RNA Communication Program**

### **Focus of this program:**

- How do cells use exRNAs to send messages to cells in distant organs?
- Do diseased cells produce different exRNAs than healthy cells, and what is the impact of these exRNAs? Can diseases like cancer spread through release of exRNAs?
- Can researchers harness the communication powers of exRNAs to turn a diseased cell into a healthy cell? (Therapeutic potential.)
- Can exRNAs be used as biomarkers to diagnose disease, monitor progression, and measure response to therapy?



### The ExRNA Communication Program is a Trans-NIH Effort

#### **ExRNA Communication Program Working Group:**

#### **Co-Chairs:**

Chris P. Austin, M.D. (NCATS) Dinah S. Singer, Ph.D. (NCI)

#### **Working Group Coordinators:**

Kevin Howcroft, Ph.D. (NCI) Patricia Labosky, Ph.D. (OD) Danilo A. Tagle, Ph.D. (NCATS)

#### Project Leaders:

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#### Program Analysts/Managers:

Tania Lombo, Ph.D. (NCATS) Dena Procaccini (NIDA)

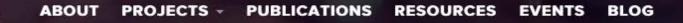
#### **Trans- NIH Members:**

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### **Program Initiatives**

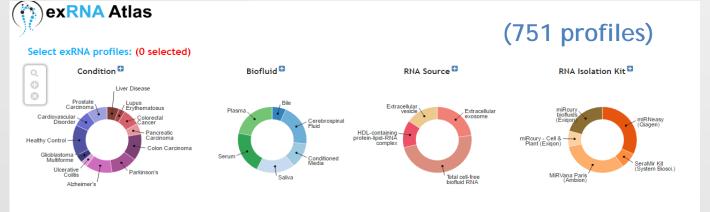
#### Defining a Comprehensive <u>Reference Profile</u> of Circulating Human Extracellular RNAs (U01)

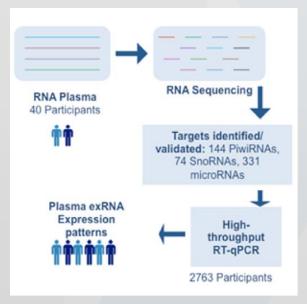
- Goal: To develop reference profiles for non-coding regulatory exRNAs from healthy human body fluids.
- Extracellular RNA Biogenesis, Biodistribution, Uptake, and Effector Function (U19)
  - Goal: To determine the biological principles guiding exRNA biogenesis, biodistribution, uptake, and effector function.
- Clinical Utility of Extracellular RNA for <u>Biomarker Development (UH2/UH3)</u>
  - Goal: To identify and quantify exRNA-based biomarkers derived from human body fluids in order to diagnose diseases, monitor disease progression, and measure response to therapy.
- Clinical Utility of Extracellular RNA for <u>Therapy Development (UH2/UH3)</u>
  - Goal: To develop and demonstrate the potential for clinical utility of exRNAs as therapeutic agents.
- Data Management and Resource Repository (DMRR)
  - Goal: To integrate the efforts of all ExRNA Communications program components and serve as a community-wide resource for exRNA standards, protocols, and data.



Exrna.org – a portal for broad dissemination of resources and data

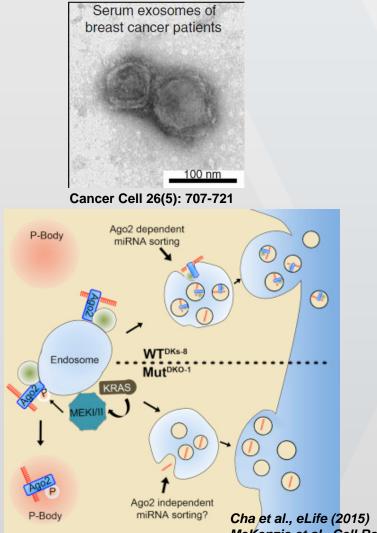
### Defining a Comprehensive Reference Profile of Circulating Human Extracellular RNAs (U01)





- Defining RNA profiles in Bile, CFS, Serum, Plasma, etc (751 profiles available)
- This represents "normal" individuals for comparison to experimental/disease/affected cohort.
- Appreciation for complexity and diversity in all sources, including plasma<sup>1</sup>:
  - Extracellular small RNAs are widely detected in the circulation in large populations
  - identified over a thousand human extracellular RNAs including microRNAs, piwi-interacting RNA (piRNA), and small nucleolar RNAs
  - Non-human RNAs detected

### Extracellular RNA Biogenesis, Biodistribution, Uptake, and Effector Function (U19)



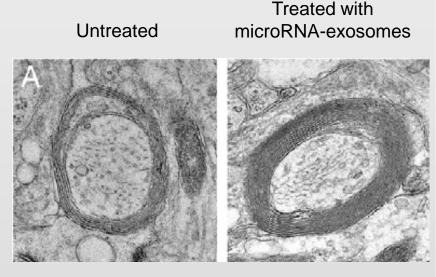
- Cancer exosomes use Dicer, TRBP, and AGO2 to process pre-miRNAs to generate mature tumor inducing miRNAs.
- Cancer exosomes mediate efficient and rapid silencing of mRNAs to reprogram a target cell transcriptome.
- **KRAS-MEK-ERK** signaling promotes Ago2 phosphorylation which inhibits Ago2-endosome association and sorting to exosomes
- Ago2 levels and phosphorylation status control secretion of candidate miRNAs in exosomes
- Indication that RNPs (like Ago2) likely very involved in chaperoning RNAs to final destinations.
- First demonstration that cell surface receptor signaling can determine loading of a specific miRNA into vesicles

Cha et al., eLife (2015) McKenzie et al., Cell Reports (2016) Calin (University of Texas) Coffey, Weaver, Patton (Vanderbilt)

### Clinical Utility of Extracellular RNA for Therapy Development (UH2/UH3):Neurological Diseases

Dr. Richard Kraig (University of Chicago) has shown microRNA-containing exosomes promote the formation of myelin in animal models of multiple sclerosis.

- Exosomes promoting myelin formation are released from immune cells stimulated with IFNγ, and are also produced by young animals and animals experiencing environmental enrichment.
- These exosomes contain microRNAs that promote oligodendrocyte differentiation and protect against inflammation.
- MicroRNA-containing exosomes promote myelin formation in the brain when administered nasally



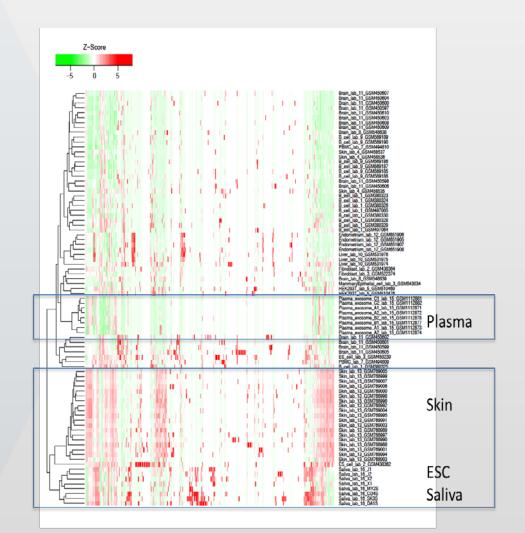
### Clinical Utility of Extracellular RNA for Biomarker Development (UH2/UH3): Gastric Cancer

Verified biomarkers for validation: 9 Long RNAs and 7 small RNAs (100 GC and 100 non-GC).

Final cohort: Saliva from 750 GC patients and 750 non-GC controls Individual biomarker validation (500 GC/ 500 non-GC)

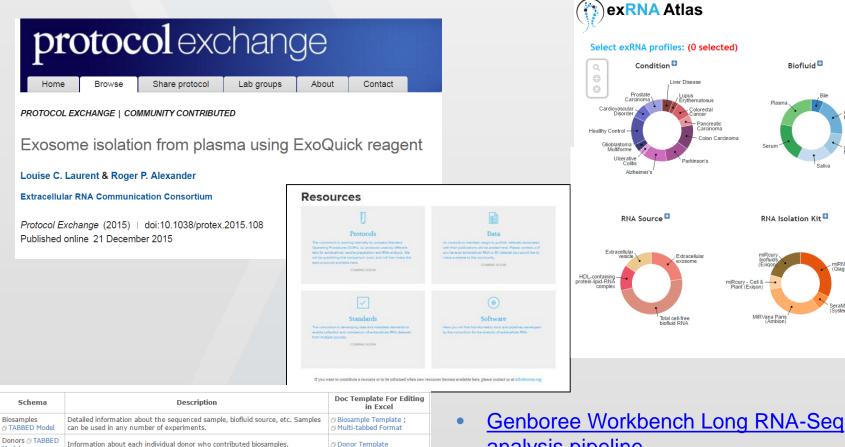
Salivary exRNA biomarker panel configuration

Next: Identification of **bacterial exRNA** biomarkers for gastric cancer in salivary



Bahn JH, Zhang Q, Li F, Chan TM, Lin X, Kim Y, Wong DT, Xiao X. The landscape of microRNA, Piwiinteracting RNA, and circular RNA in human saliva. Clin Chem. 2015;61(1): 221-30. PMCID: 4332885.

### **Data Management and Resource Repository (DMRR)**



Study Template

Experiment Template

Submission Template

Analysis Template

Run Template

Model Studies 🗇 TABBED

Model

Analyses

Model

Experiments

□ TABBED Model

TABBED Model Submissions

TABBED Model Runs @ TABBED

purposes.

groups together one or more runs.

An analysis contains secondary analysis results.

Information about PI / submitter associated with submission.

A run contains sequencing reads submitted in data files.

A study groups together experiments or analyses for public data release

An experiment contains instrument and library preparation information and

- analysis pipeline exceRpt - Genboree Workbench Small RNA-
- seq analysis pipeline

Biofluid

erebrospinal Iuid

Qiagen)

eraMir Kit System Biosci

### **Some Consortium Wide Accomplishments**

- 197 total publications in 50 different scientific journals, demonstrating broad reach of the program into various biomedical specialties.
- Generation of highly standardized protocols for vesicle isolation and RNA purification from body fluids, body fluid collection and processing, etc.
- Revision of Gene Ontology terms (GO terms) relevant for exRNA-containing particles.
- Developed standards and standard formats, tools and applications, for RNA-Seq data and metadata standards, which is essential to data harmonization
- Pls are leading a new Gordon Conference and a Keystone Meeting.
- Developing imaging tools to visualize and track vesicles.
- Developing a set of cell lines with mutations in all the known components of vesicle biogenesis.



### **Remaining Challenges**

- ExRNA communication mechanisms are in early discovery stage
- The consortium has gelled and is working together to address some of the cross-cutting fundamental discovery goals of the program.
- All therapy (8) and biomarker (10) projects are moving forward
  - oligodendrocyte differentiation
  - siRNA targeting of mutant Huntingtin mRNA
  - edible plant-derived nano-vector delivery
  - biomarkers for various cancers, cardiovascular risk, placental dysfunction, CNS disease
- However, successful therapeutics should have a foundation of defined biological principles behind them. Some therapeutic goals may be dependent on determining underlying biology – which takes more time.
- Much of this foundational work is hypothesis-generating and broadly enabling, but are there are opportunities for additional focused investments to reveal new paradigms within another 4-5 years?

