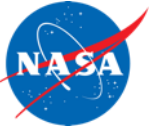


# Circulating miRNA Signature Predicts Health Risks Associated with Radiation and Microgravity

National Aeronautics and  
Space Administration



**Afshin Beheshti, PhD**  
**Bioinformatician at GeneLab**  
**Principal Investigator**  
**Space Biosciences Division, KBRWyle**  
**NASA Ames Research Center, Moffett Field, CA**

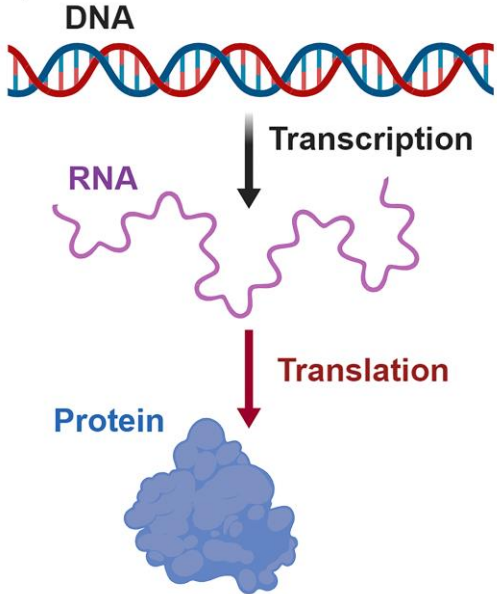
**Adjunct Assistant Professor at Department of Medicine**  
**Rutgers Robert Wood Johnson Medical School**

**Visiting Researcher at Broad Institute**  
**Cambridge, MA**

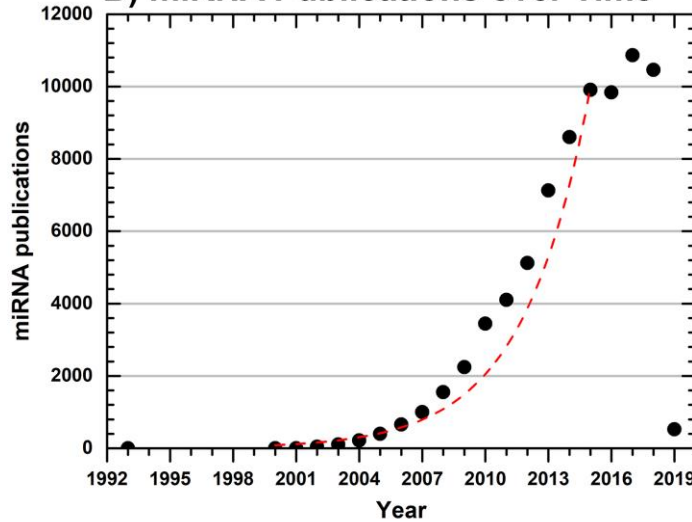
**[afshin.beheshti@nasa.gov](mailto:afshin.beheshti@nasa.gov)**  
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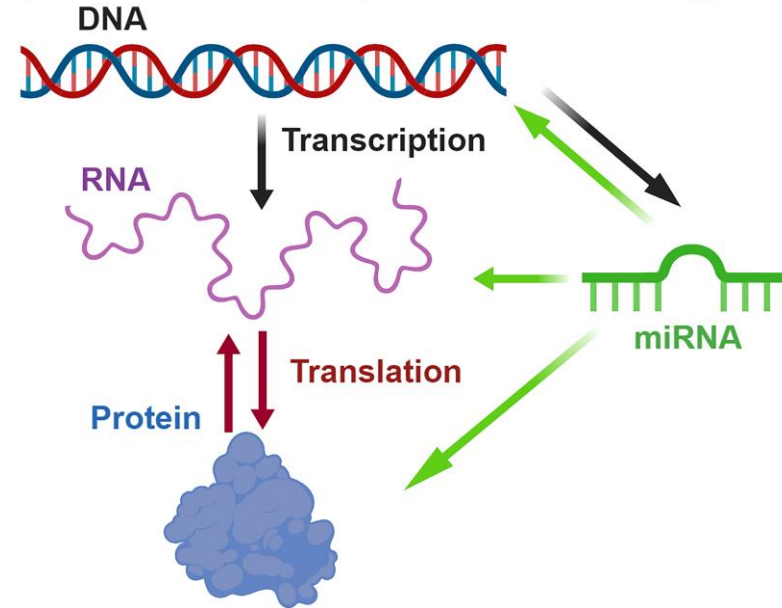
## A) Classical View of Molecular Biology



## B) miRNA Publications over Time



## C) New Understanding of Molecular Biology



- A single miRNA has been estimated to regulate up to 500 mRNAs.
- miRNAs are ~22nt
- **Due to the size and stability of the miRNAs, it can float freely in the blood.**
- miRNAs are now known to be involved in all aspects of diseases.
- miRNA are not only found in mammals, but everything else living: plants, microbes, fish, C. Elegans, fruit flies, insects, etc...
- miRNAs play a big role in radiation response (which also relates to space radiation).

Saliva	Breast Milk	Venous Blood	Cerebrospinal Fluid	Tears	Peritoneal Fluid	Semen	Vaginal Secretion	Menstrual Blood	Plasma		
miR-26a <sup>[c]</sup> miR-96 <sup>[c]</sup> miR-135b <sup>[c]</sup> miR-141 <sup>[c]</sup> miR-145 <sup>[c]</sup> miR-182 <sup>[c]</sup>	miR-10a <sup>[c]</sup> miR-28-5p <sup>[c]</sup> miR-150 <sup>[c]</sup> miR-193b <sup>[c]</sup> miR-217 <sup>[c]</sup> miR-518c <sup>[c]</sup> miR-924 <sup>[c]</sup>	miR-16 <sup>[a,b]</sup> miR-20a <sup>[b]</sup> miR-106a <sup>[b]</sup> miR-126 <sup>[d]</sup> miR-150 <sup>[d]</sup> miR-185 <sup>[b]</sup> miR-451 <sup>[a,b,d]</sup> miR-451a <sup>[f]</sup>	miR-577 <sup>[c]</sup>	miR-637 <sup>[c]</sup>	miR-29b-1 <sup>[c]</sup> miR-129 <sup>[c]</sup>	miR-10a <sup>[b]</sup> miR-10b <sup>[a,b]</sup> miR-17 <sup>[c]</sup> miR-29b-2 <sup>[c]</sup> miR-135a <sup>[b]</sup>	miR-144 <sup>[b]</sup> miR-124a <sup>[a]</sup> miR-372 <sup>[c]</sup> miR-617 <sup>[b]</sup>	miR-223 <sup>[c]</sup> miR-583 <sup>[c]</sup>	miR-518f <sup>[c]</sup> miR-135a <sup>[c]</sup> miR-139-3p <sup>[c]</sup> miR-182 <sup>[c]</sup> miR-224 <sup>[c]</sup> miR-299-5p <sup>[c]</sup>	miR-508-5p <sup>[c]</sup> miR-644 <sup>[c]</sup> miR-891a <sup>[b]</sup> miR-943 <sup>[b]</sup> miR-185 <sup>[b]</sup> miR-412 <sup>[a]</sup> miR-451 <sup>[a]</sup> miR-330-5p <sup>[c]</sup> miR-369-3p <sup>[c]</sup> miR-373 <sup>[c]</sup> miR-483-3p <sup>[c]</sup> miR-508-3p <sup>[c]</sup>	miR-519d <sup>[c]</sup> miR-551b <sup>[c]</sup> miR-801 <sup>[c]</sup>

Silva, S.S., et al., *Forensic miRNA: potential biomarker for body fluids?* Forensic Sci Int Genet, 2015, 14: p. 1-10.

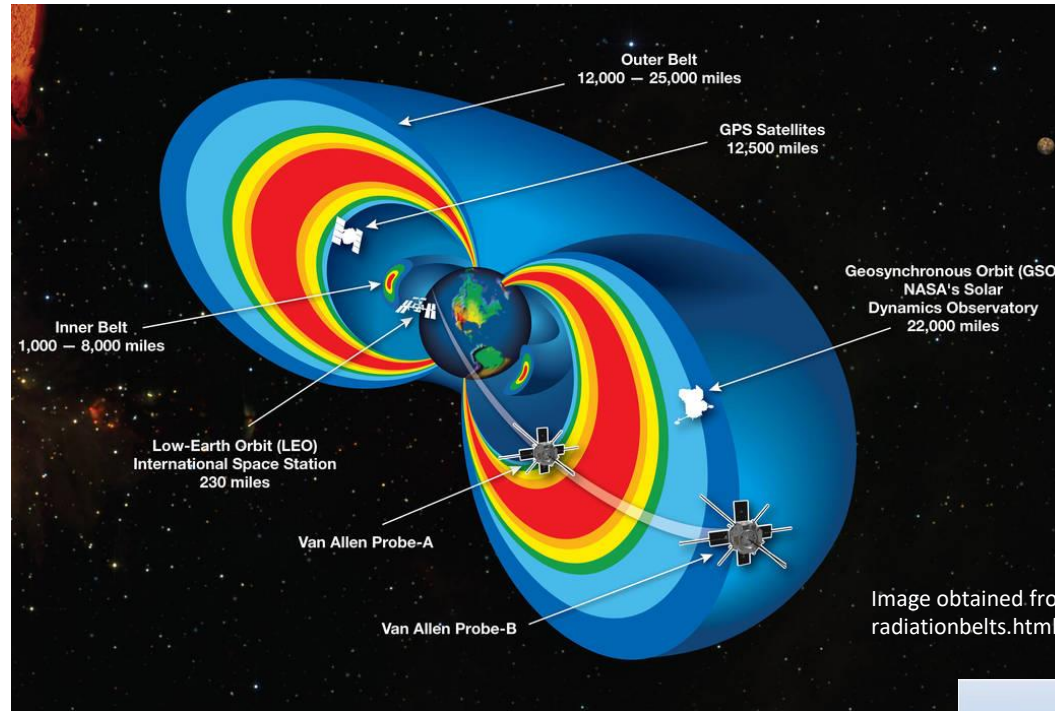
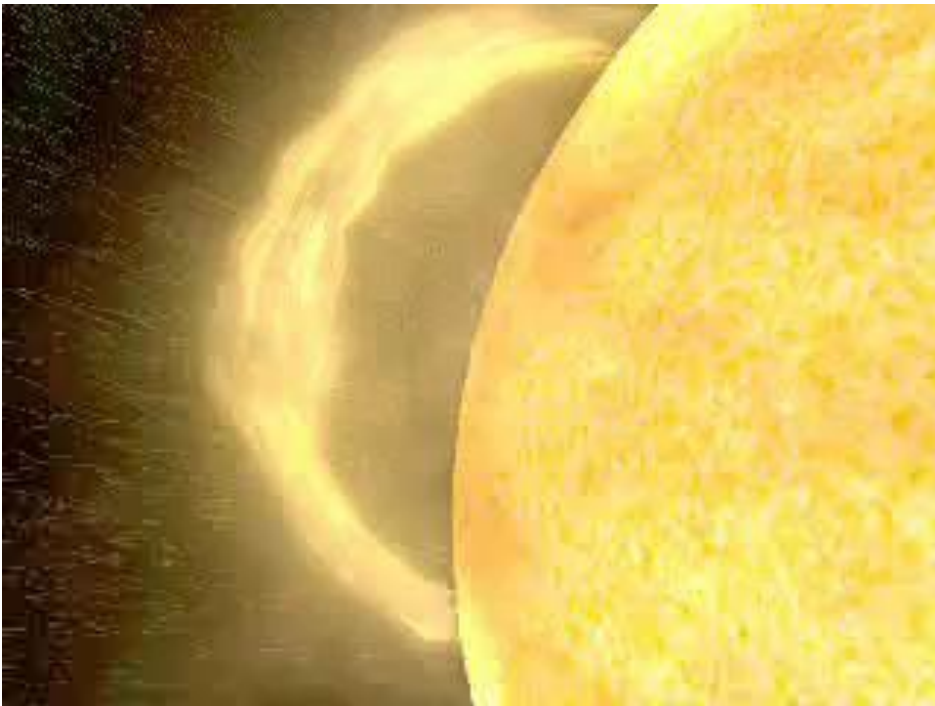


Image obtained from [radiationbelts.html](http://radiationbelts.html)

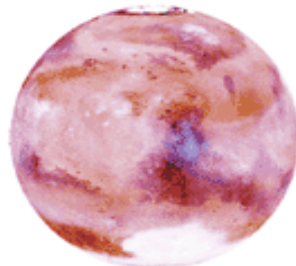
MILLIREM:

CHEST X-RAY	8 to 50
AVG. YEARLY RADON DOSE	200
U.S. AVG. YEARLY DOSE	350
PET SCAN	1,000
1 YEAR IN KERALA, INDIA	1,300
U.S. NUCLEAR WORKER LIMIT PER YEAR	5,000
APOLLO 14 (9 DAYS)	1,140
SHUTTLE 41-C (18 DAYS)	5,600
SKYLAB 4 (84 DAYS)	17,800
MARS MISSION TOTAL	130,000

**2½ Years, 2,600 X-Rays**

Americans on average absorb the radiation equivalent of at least 7 chest X-rays each year.

Space missions, outside of Earth's protective atmosphere and magnetic field, expose astronauts to many times more.



NASA

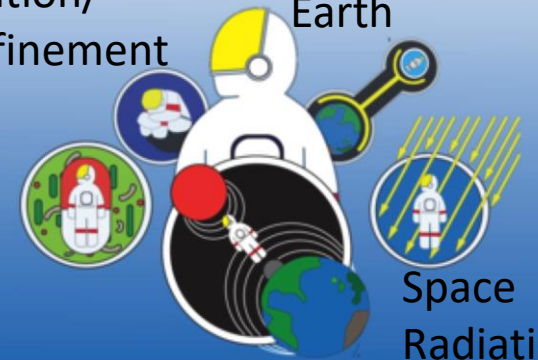
Source: Brookhaven National Laboratory, U.S. Department of Energy



Isolation/  
Confinement

Distance from  
Earth

Hostile/closed  
environments



Space  
Radiation

Gravity Fields



## FEMALE ASTRONAUT



Women suffer less from hearing loss with advancing age, and do not display a bias towards loss of hearing in the left ear

Women demonstrate a slight bias towards accuracy versus speed in response to an alertness test

Women mount more potent immune responses

Struvite kidney stones more common in women

Female astronauts, (to date) do not exhibit clinically significant visual impairment

Female astronauts are more susceptible to orthostatic intolerance

Urinary tract infections are more common in female astronauts

Large individual variability to muscle and bone loss in women

Health effect observed on Earth

## MALE ASTRONAUT



Men suffer more from hearing loss with advancing age, and display a bias towards loss of hearing in the left ear

Men demonstrate a slight bias towards speed versus accuracy in response to an alertness test

Men mount less potent immune responses

Calcium oxalate kidney stones more common in men

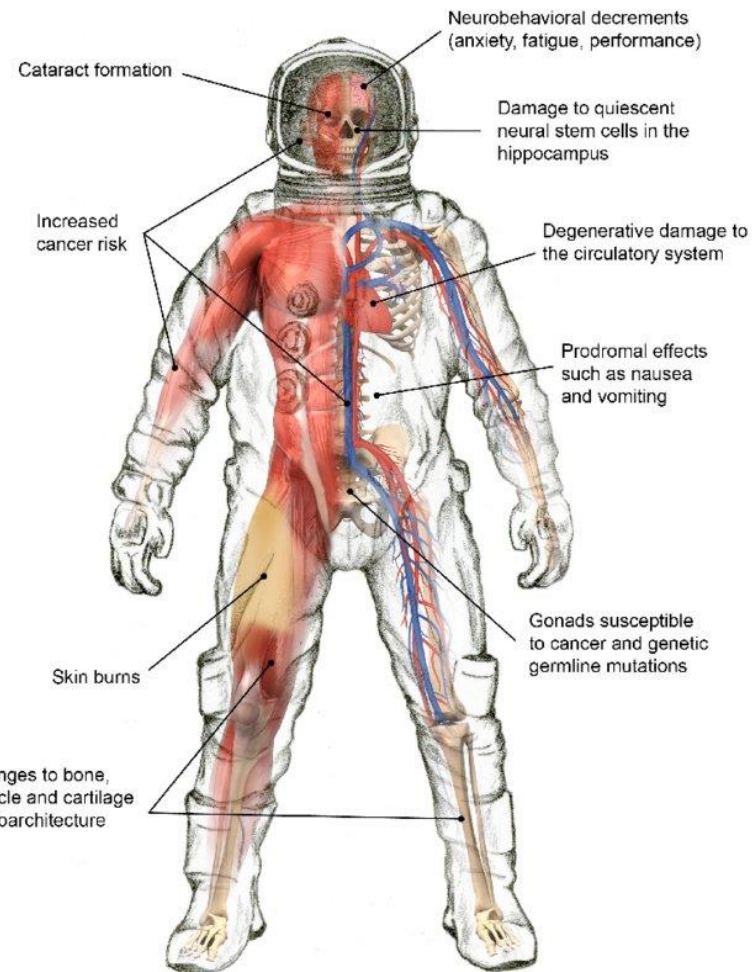
Some male astronauts exhibit clinically significant visual impairment

Male astronauts less susceptible to orthostatic intolerance

Urinary tract infections less common in male astronauts

Large individual variability to muscle and bone loss in men

Health effect observed in space



Select health effects due to space radiation exposures.

From: J. Chancellor et al., Space Radiation: The Number One Risk to Astronaut Health beyond Low Earth Orbit. *Life*, 4(3), 491-510;

## Systems Biology View of miRNAs

RESEARCH ARTICLE

A microRNA signature and TGF- $\beta$ 1 response were identified as the key master regulators for spaceflight response

Afshin Beheshti<sup>1\*</sup>, Shayoni Ray<sup>2\*</sup>, Homer Fogle<sup>1</sup>, Daniel Berrios<sup>2</sup>, Sylvain V. Costes<sup>3\*</sup>

**1** WYLE, NASA Ames Research Center, Moffett Field, California, United States of America, **2** USRA, NASA Ames Research Center, Moffett Field, California, United States of America, **3** NASA Ames Research Center, Space Biosciences Division, Moffett Field, California, United States of America

\* These authors contributed equally to this work.

\* afshin.beheshti@nasa.gov (AB); sylvain.v.costes@nasa.gov (SVC)



### Abstract

Translating fundamental biological discoveries from NASA Space Biology program into health risk from space flights has been an ongoing challenge. We propose to use NASA GeneLab database to gain new knowledge on potential systemic responses to space. Unbiased systems biology analysis of transcriptomic data from seven different rodent datasets reveals for the first time the existence of potential "master regulators" coordinating a systemic response to microgravity and/or space radiation with TGF- $\beta$ 1 being the most common regulator. We hypothesized the space environment leads to the release of biomolecules circulating inside the blood stream. Through datamining we identified 13 candidate microRNAs (miRNA) which are common in all studies and directly interact with TGF- $\beta$ 1 that can be potential circulating factors impacting space biology. This study exemplifies the utility of the

### OPEN ACCESS

**Citation:** Beheshti A, Ray S, Fogle H, Berrios D, Costes SV (2018) A microRNA signature and TGF- $\beta$ 1 response were identified as the key master regulators for spaceflight response. PLoS ONE 13 (7): e0199621. <https://doi.org/10.1371/journal.pone.0199621>

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**Accepted:** May 3, 2018

**Published:** July 25, 2018



### Article

## GeneLab Database Analyses Suggest Long-Term Impact of Space Radiation on the Cardiovascular System by the Activation of *FYN* Through Reactive Oxygen Species

Afshin Beheshti<sup>1,\*</sup>, J. Tyson McDonald<sup>2</sup>, Jack Miller<sup>3</sup>, Peter Grabham<sup>4</sup> and Sylvain V. Costes<sup>5,\*</sup>

**1** WYLE Labs, NASA Ames Research Center, Moffett Field CA 94035, USA

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**4** Center for Radiological Research, Columbia University, New York, NY 10032, USA; pwg2@cumc.columbia.edu

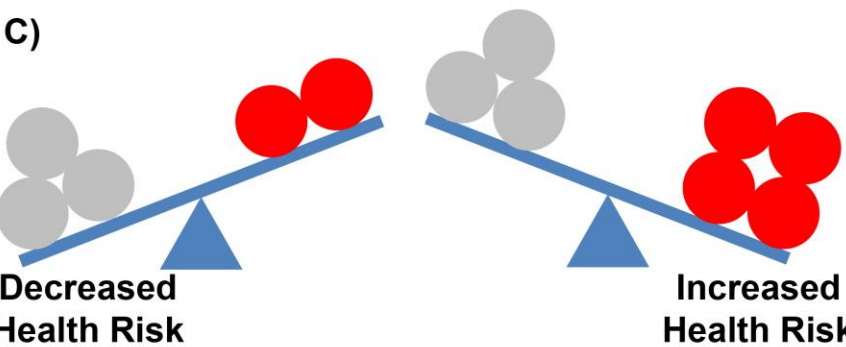
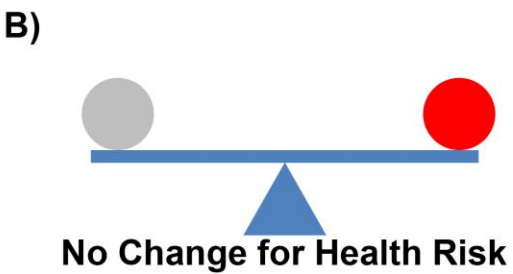
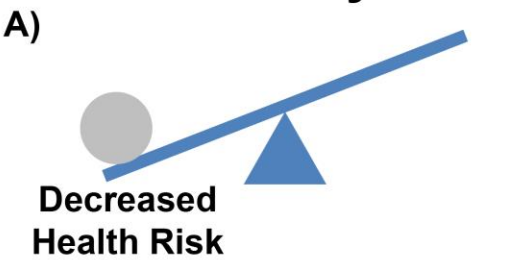
**5** NASA Ames Research Center, Space Biosciences Division, Moffett Field, CA 94035, USA

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Tel.: +1-650-604-5343 (S.V.C.)

Received: 15 January 2019; Accepted: 30 January 2019; Published: 3 February 2019

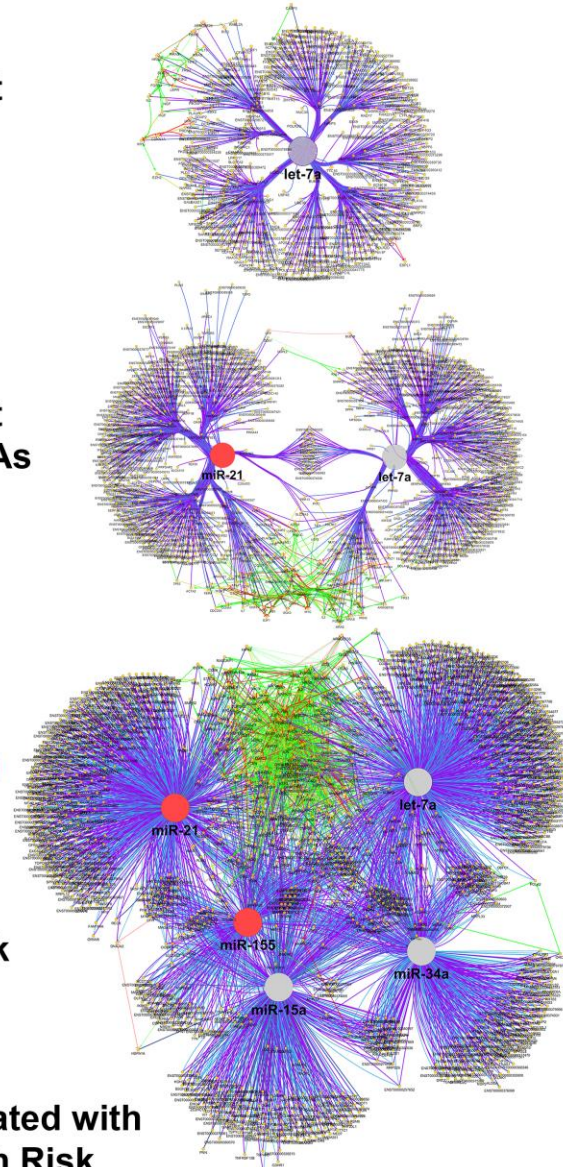
Received: 15 January 2019; Accepted: 30 January 2019; Published: 3 February 2019



**Systems Biology Approach:** Looking at how the most important miRNAs impact the entire system

● miRNAs Associated with Decreased Health Risk

● miRNAs Associated with Increased Health Risk



RESEARCH ARTICLE

## A Circulating microRNA Signature Predicts Age-Based Development of Lymphoma

Afshin Beheshti<sup>1</sup>, Charles Vanderburg<sup>2</sup>, J. Tyson McDonald<sup>3</sup>, Charushella Ramkumar<sup>4</sup>, Tatenda Kadungure<sup>4</sup>, Hong Zhang<sup>4</sup>, Ronald B. Gartenhaus<sup>5</sup>, Andrew M. Evens<sup>1\*</sup>

**1** Division of Hematology/Oncology, Molecular Oncology Research Institute, Tufts Medical Center, Boston, Massachusetts, United States of America, **2** Harvard NeuroDiscovery Center, Massachusetts General Hospital, Boston, Massachusetts, United States of America, **3** Cancer Research Center, Hampton University, Hampton, Virginia, United States of America, **4** Department of Cell Biology and Development, University of Massachusetts Medical School, Worcester, Massachusetts, United States of America, **5** Marlene & Stewart Greenbaum Cancer Center, Department of Medicine, University of Maryland, Baltimore, Maryland, United States of America

\*AEvens@tuftsmedicalcenter.org

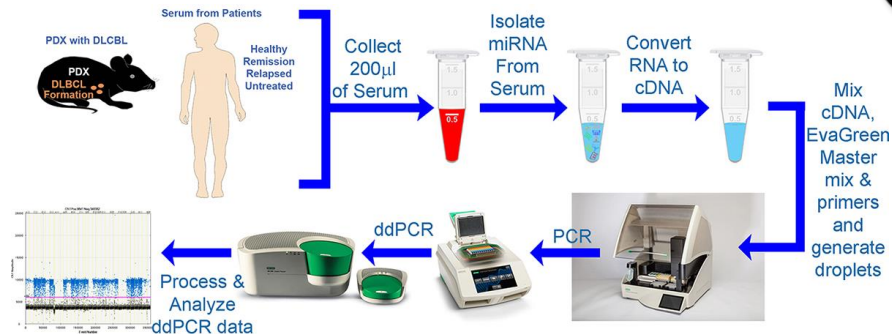
### Abstract

Extensive epidemiological data have demonstrated an exponential rise in the incidence of non-Hodgkin lymphoma (NHL) that is associated with increasing age. The molecular etiology of this remains largely unknown, which impacts the effectiveness of treatment for patients. We proposed that age-dependent circulating microRNA (miRNA) signatures in the host influence diffuse large B cell lymphoma (DLBCL) development. Our objective was to examine tumor development in an age-based DLBCL system using an inventive systems biology approach. We harnessed a novel murine model of spontaneous DLBCL initiation

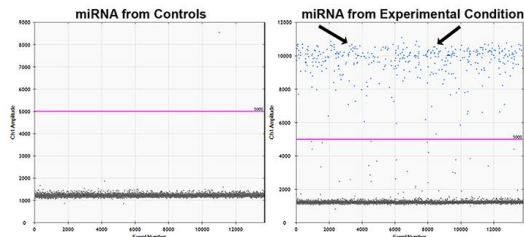


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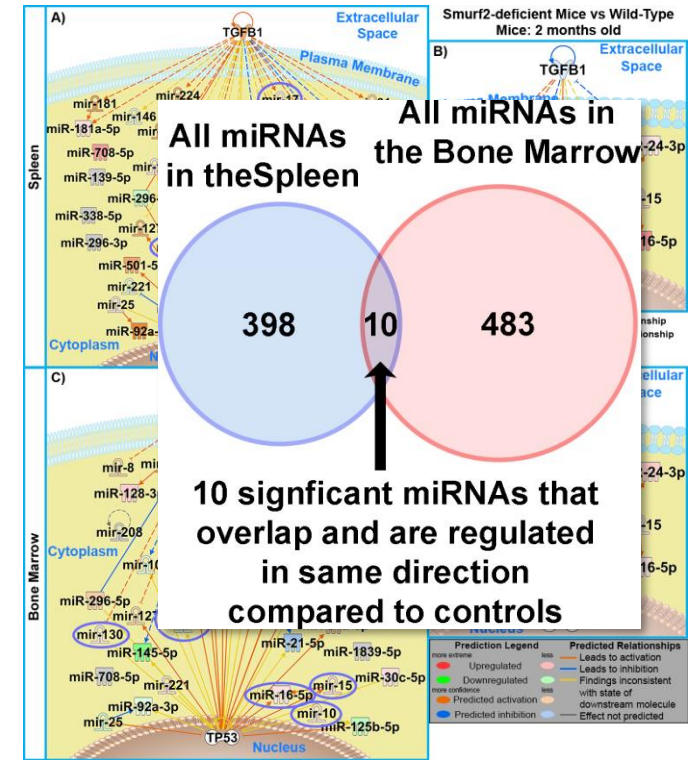
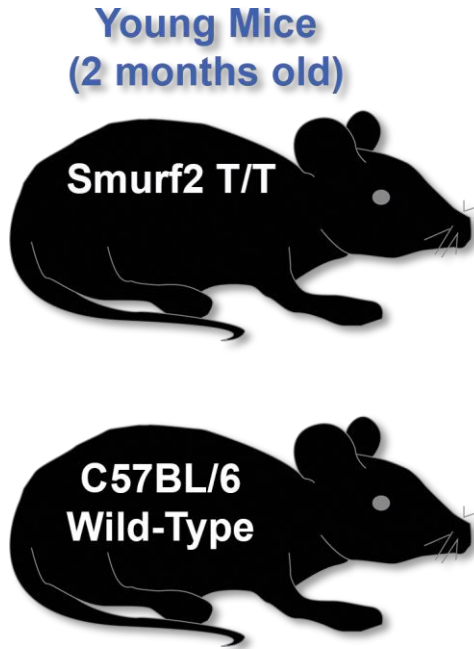
**Citation:** Beheshti A, Vanderburg C, McDonald JT, Ramkumar C, Kadungure T, Zhang H, et al. (2017) A Circulating microRNA Signature Predicts Age-Based Development of Lymphoma. PLOS ONE 12(1): e0170521. doi:10.1371/journal.pone.0170521



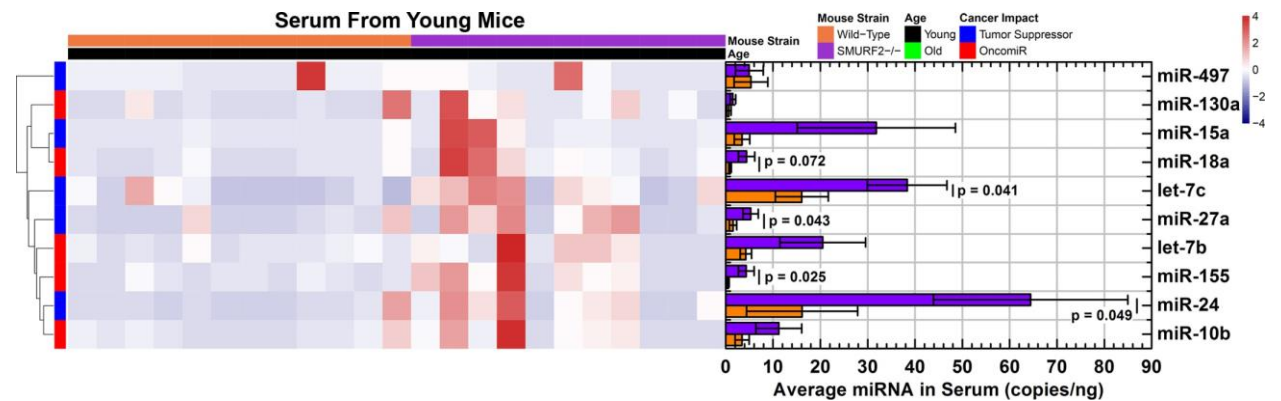
Through ddPCR we are able to get exact counts of circulating miRNA in the serum

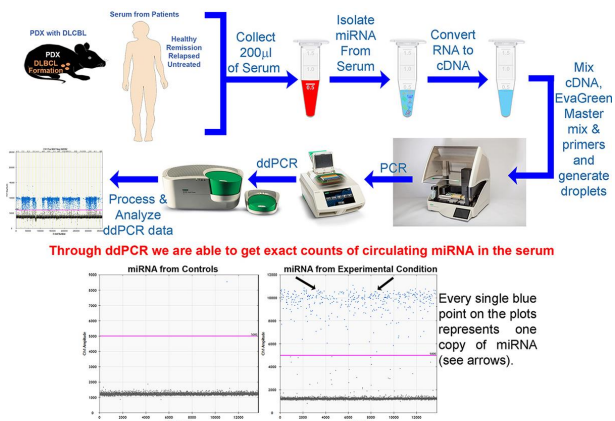
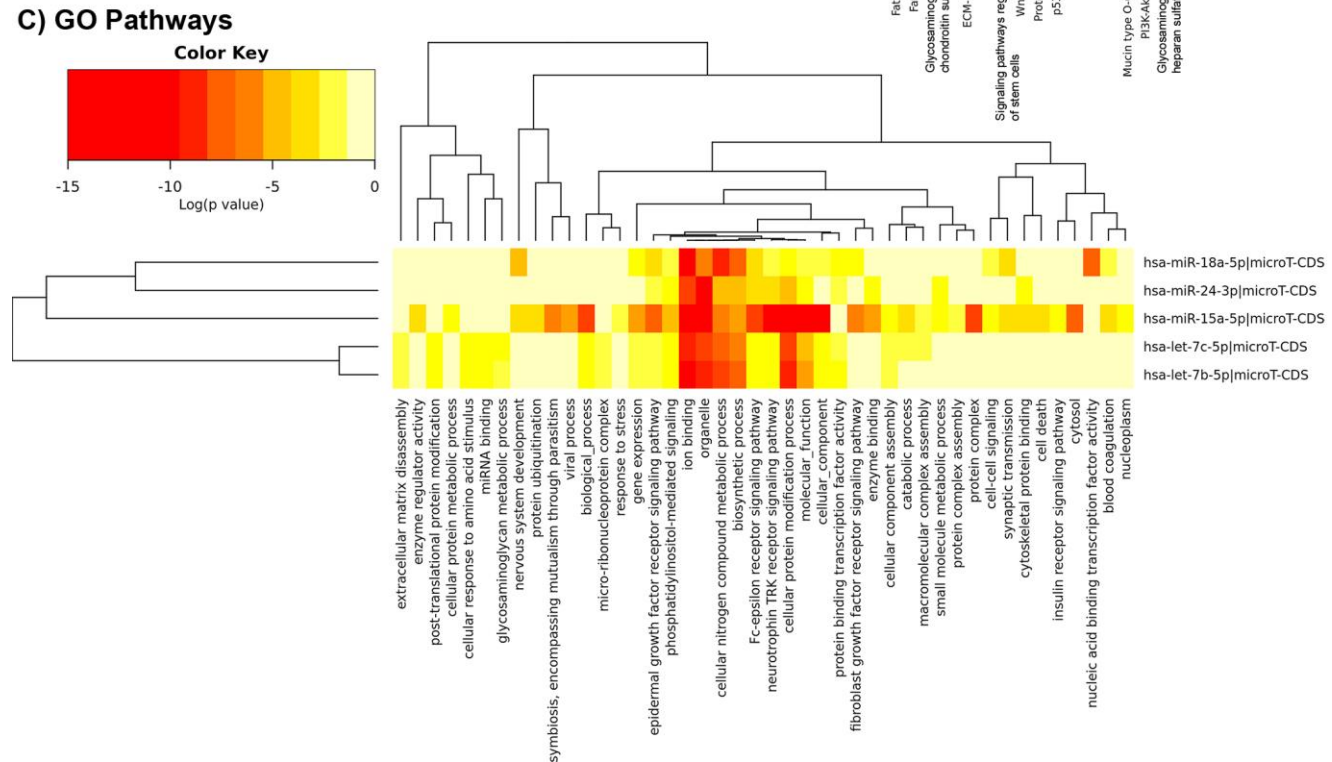
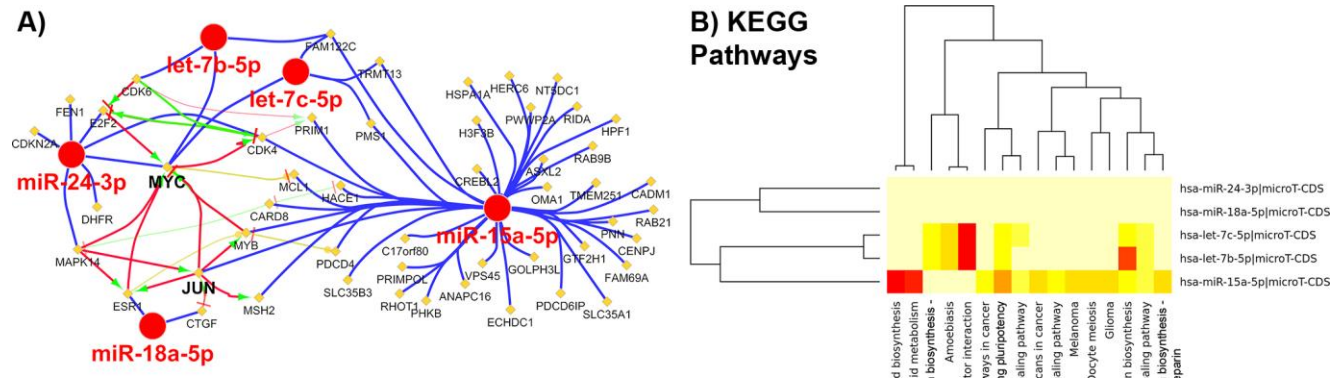
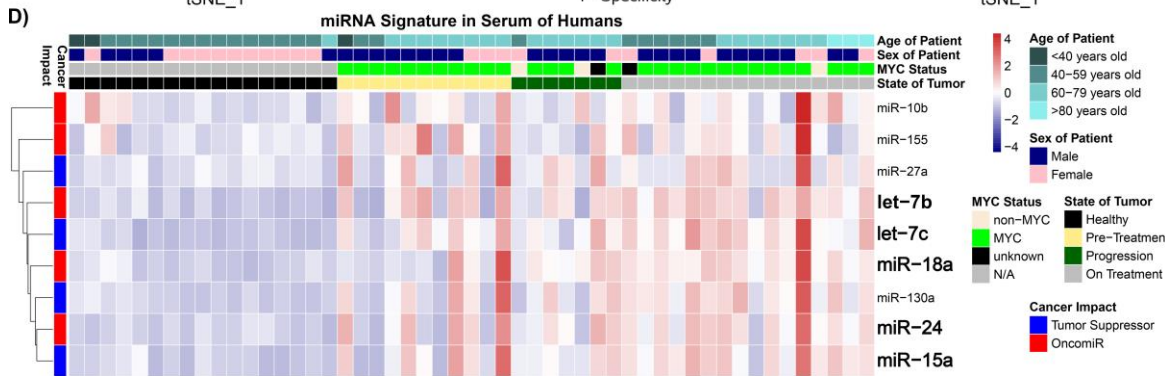
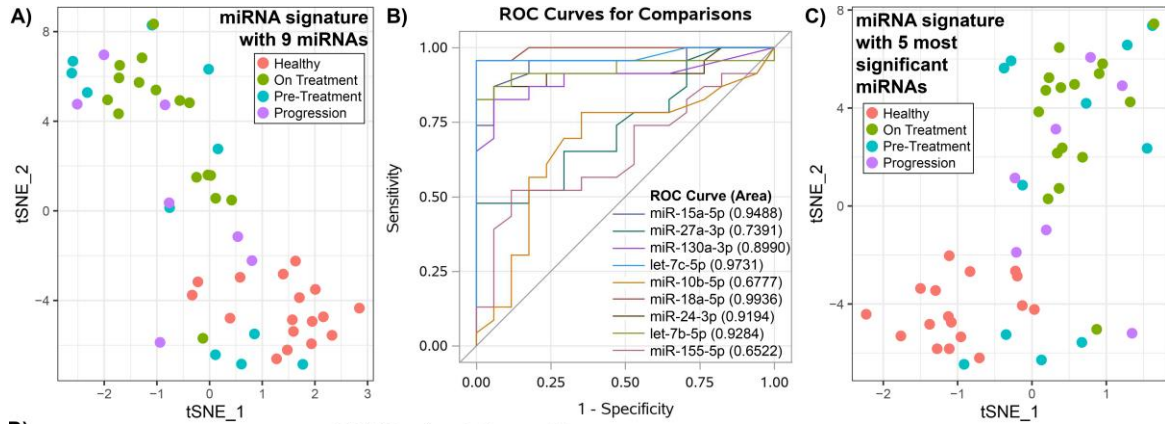


Every single blue point on the plots represents one copy of miRNA (see arrows).



10 significant miRNAs that overlap and are regulated in same direction compared to controls





RESEARCH ARTICLE

A microRNA signature and TGF- $\beta$ 1 response were identified as the key master regulators for spaceflight response

Afshin Beheshti<sup>1\*</sup>, Shayoni Ray<sup>2\*</sup>, Homer Fogle<sup>1</sup>, Daniel Berrios<sup>2</sup>, Sylvain V. Costes<sup>3\*</sup>

**1** WYLE, NASA Ames Research Center, Moffett Field, California, United States of America, **2** USRA, NASA Ames Research Center, Moffett Field, California, United States of America, **3** NASA Ames Research Center, Space Biosciences Division, Moffett Field, California, United States of America

\* These authors contributed equally to this work.  
\* afshin.beheshti@nasa.gov (AB); shayoni.v.costes@nasa.gov (SVC)



Abstract

Translating fundamental biological discoveries from NASA Space Biology program into health risk from space flights has been an ongoing challenge. We propose to use NASA GeneLab database to gain new knowledge on potential systemic responses to space. Unbiased systems biology analysis of transcriptomic data from seven different rodent datasets reveals for the first time the existence of potential "master regulators" coordinating a systemic response to microgravity and/or space radiation with TGF- $\beta$ 1 being the most common regulator. We hypothesized the space environment leads to the release of biomolecules circulating inside the blood stream. Through datamining we identified 13 candidate microRNAs (miRNA) which are common in all studies and directly interact with TGF- $\beta$ 1 that can be potential circulating factors impacting space biology. This study exemplifies the utility of the GeneLab data repository to aid in the process of performing novel hypothesis-based research.

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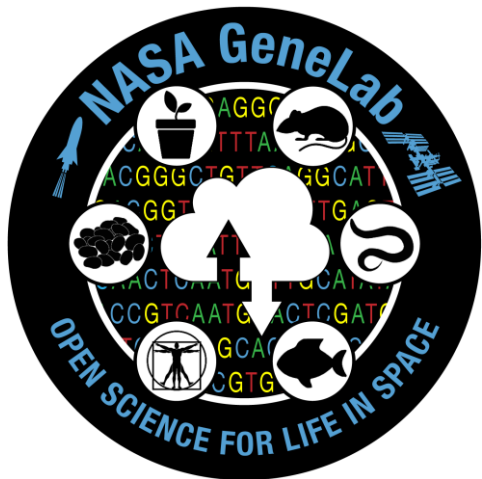
**Citation:** Beheshti A, Ray S, Fogle H, Berrios D, Costes SV (2018) A microRNA signature and TGF- $\beta$ 1 response were identified as the key master regulators for spaceflight response. PLoS ONE 13 (7): e0199621. <https://doi.org/10.1371/journal.pone.0199621>

**Editor:** Andre van Wijnen, University of Massachusetts Medical School, UNITED STATES

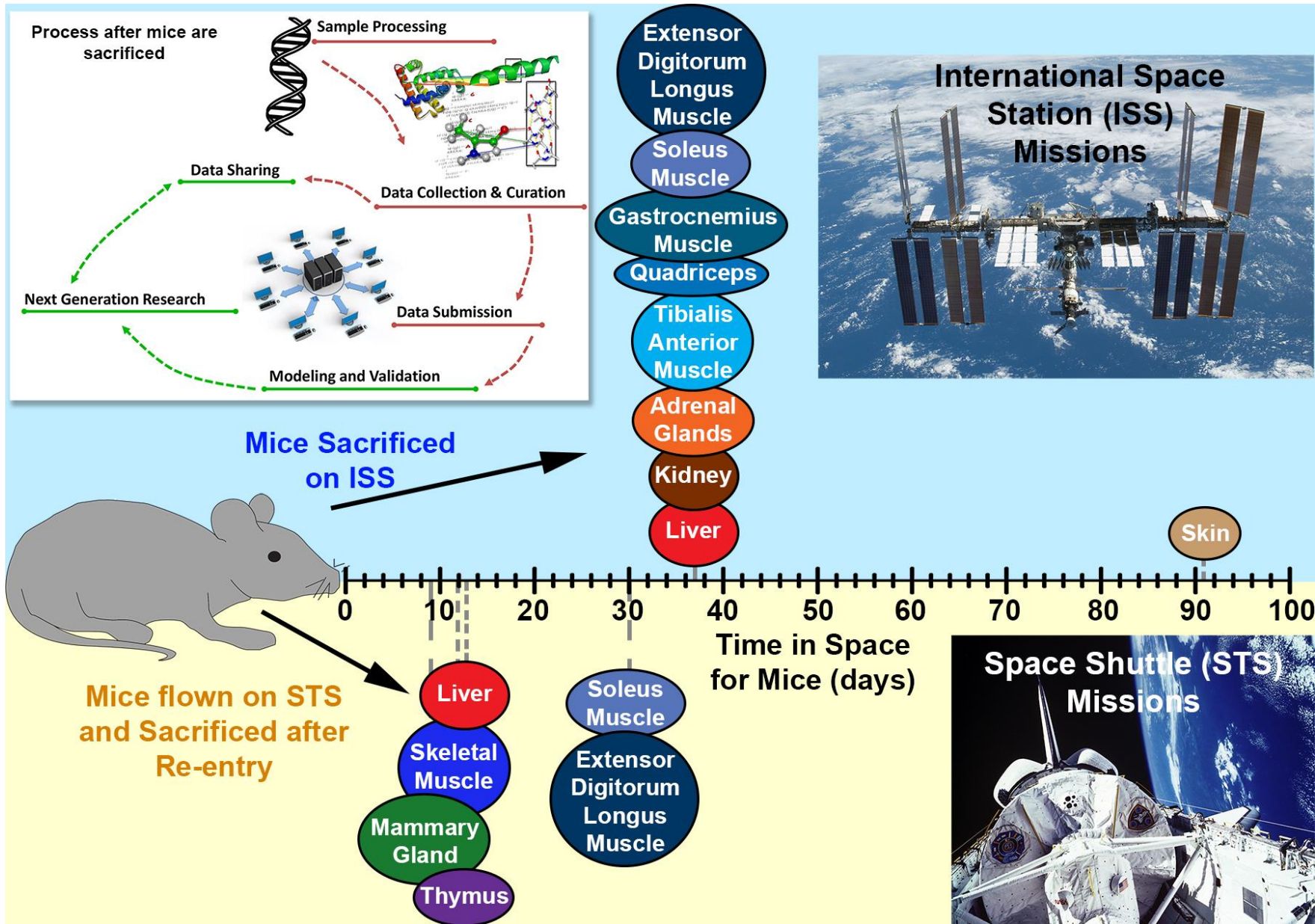
**Received:** March 6, 2018

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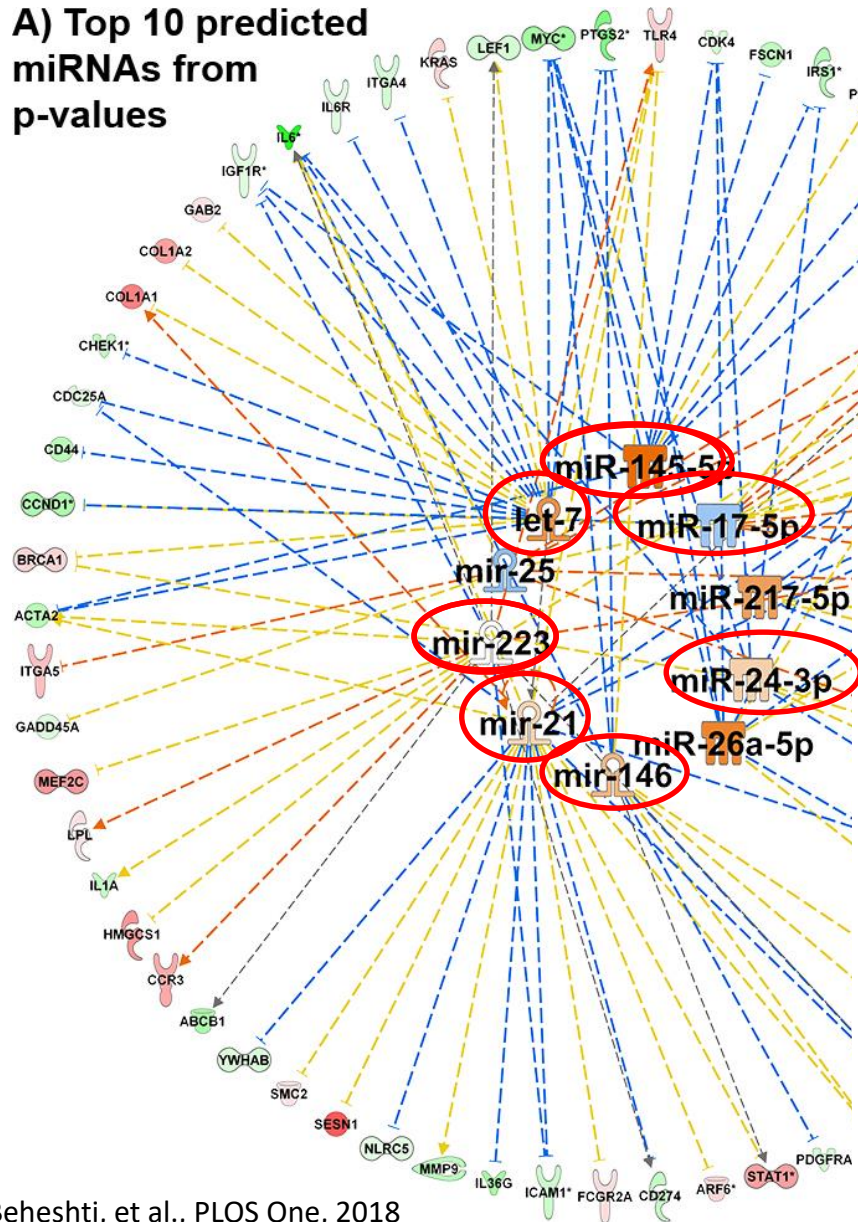


<https://genelab.nasa.gov/>

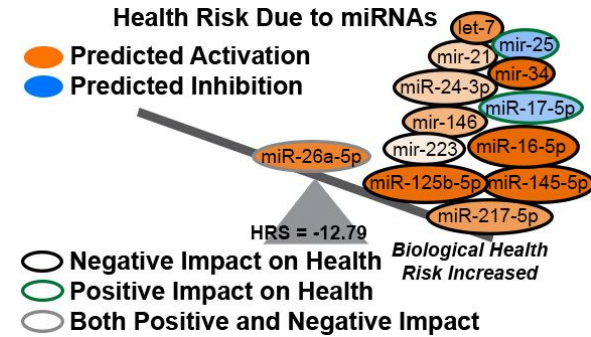
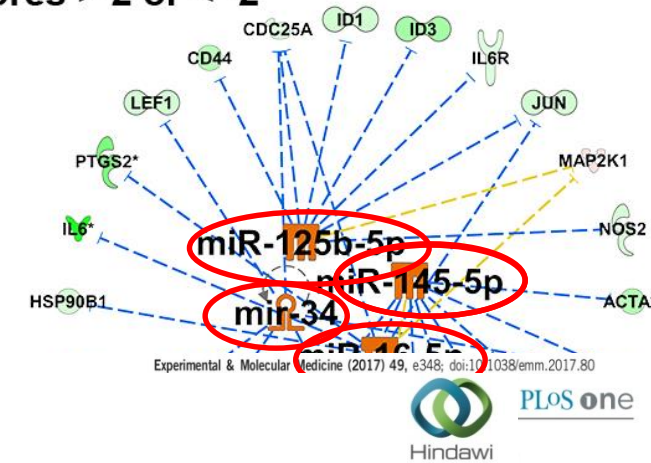




## A) Top 10 predicted miRNAs from p-values



## B) All miRNAs with Z-scores > 2 or < -2



## C) Research Article

### Integration Analysis of MicroRNA and mRNA Expression Profiles in Human Peripheral Blood Lymphocytes Cultured in Modeled Microgravity

C. Girardi,<sup>1</sup> C. De Pittà,<sup>1</sup> S. Casara,<sup>1</sup> E. Calura,<sup>1</sup> C. Romualdi,<sup>1</sup> L. Celotti,<sup>1,2</sup> and M. Mognato<sup>1</sup>

<sup>1</sup> Dipartimento di Biologia, Università degli Studi di Padova, Via U. Bassi 58/B, 35131 Padova, Italy

<sup>2</sup> Laboratori Nazionali di Legnaro, INFN, Viale dell'Università 2, Legnaro, 35020 Padova, Italy

Correspondence should be addressed to L. Celotti; lucia.celotti@unipd.it and M. Mognato; maddalena.mognato@unipd.it

Received 16 April 2014; Revised 22 May 2014; Accepted 22 May 2014; Published 23 June 2014

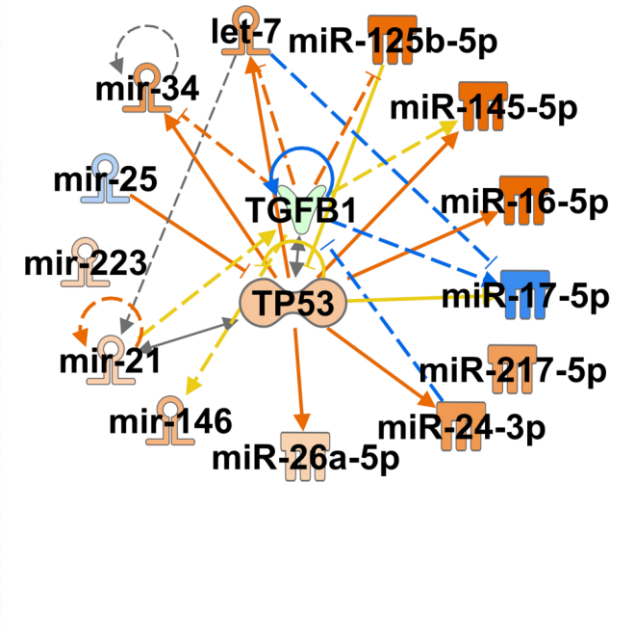
Academic Editor: Mariano Bizzarri

Copyright © 2014 C. Girardi et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

We analyzed miRNA and mRNA expression profiles in human peripheral blood lymphocytes (PBLs) incubated in microgravity condition, simulated by a ground-based rotating wall vessel (RWV) bioreactor. Our results show that 42 miRNAs were differentially expressed in MMG-incubated PBLs compared with 1g incubated ones. Among these, miR-9-5p, miR-9-3p, miR-155-5p, miR-150-3p, and miR-378-3p were the most dysregulated. To improve the detection of functional miRNA-mRNA pairs, we performed gene expression profiles on the same samples assayed for miRNA profiling and we integrated miRNA and mRNA expression data. The functional classification of miRNA-correlated genes evidenced significant enrichment in the biological processes of immune/inflammatory response, signal transduction, regulation of response to stress, regulation of programmed cell death, and regulation of cell proliferation. We identified the correlation of miR-9-3p, miR-155-5p, miR-150-3p, and miR-378-3p expression with that of genes involved in immune/inflammatory response (e.g., IFNG and IL17F), apoptosis (e.g., PDCD4 and PTEN), and cell proliferation (e.g., NKX3-1 and GADD45A). Experimental assays of cell viability and apoptosis induction validated the results obtained by bioinformatics analyses demonstrating that in human PBLs the exposure to reduced gravitational force increases the frequency of apoptosis and decreases cell proliferation.

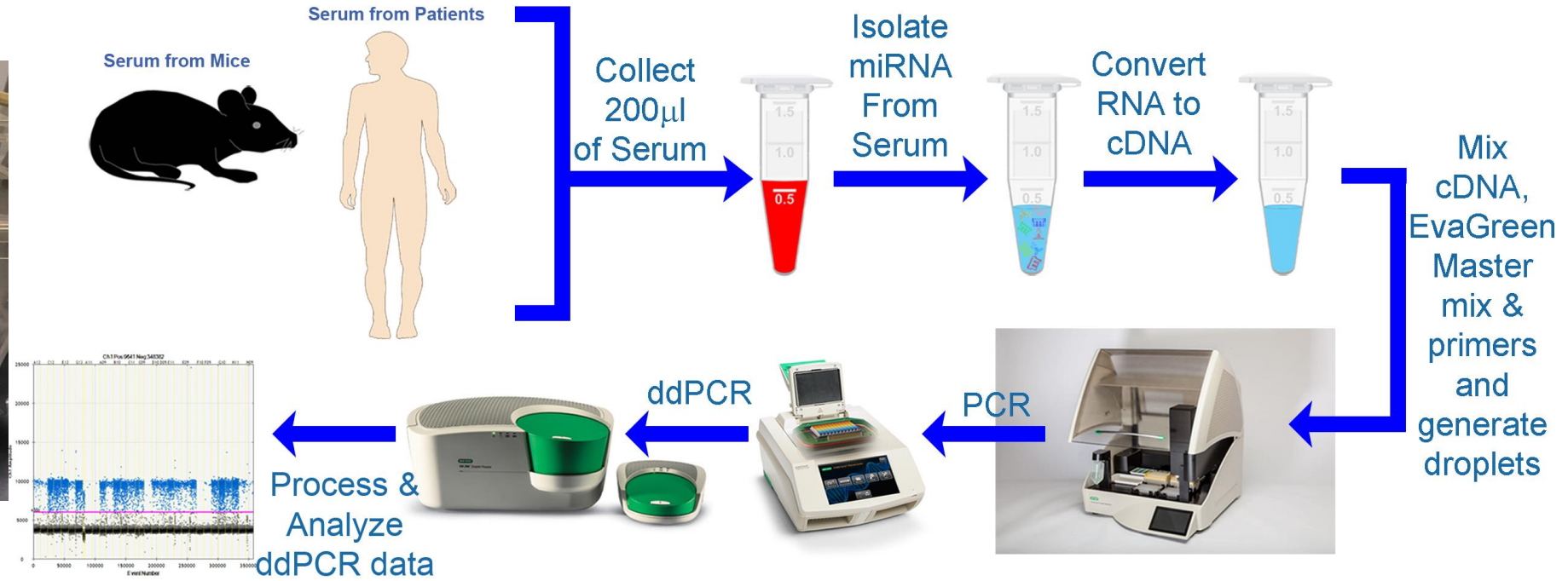
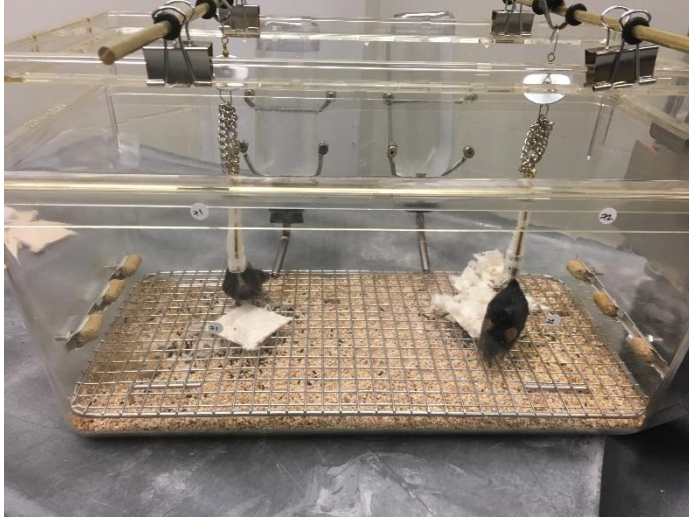
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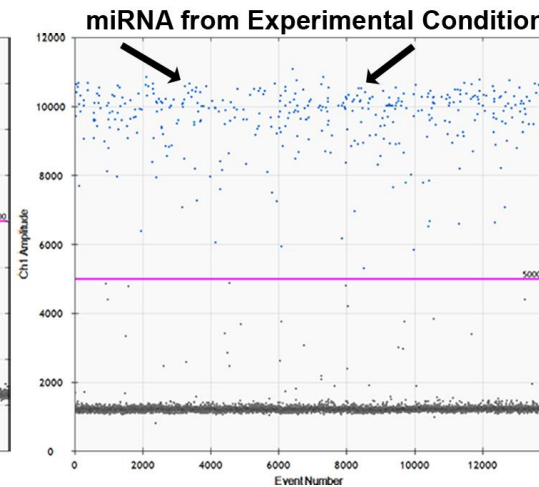
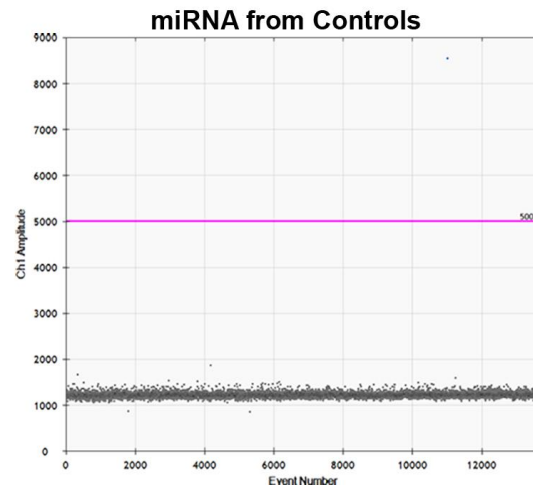


# Technique to Quantify miRNAs

## Hindlimb Unloading



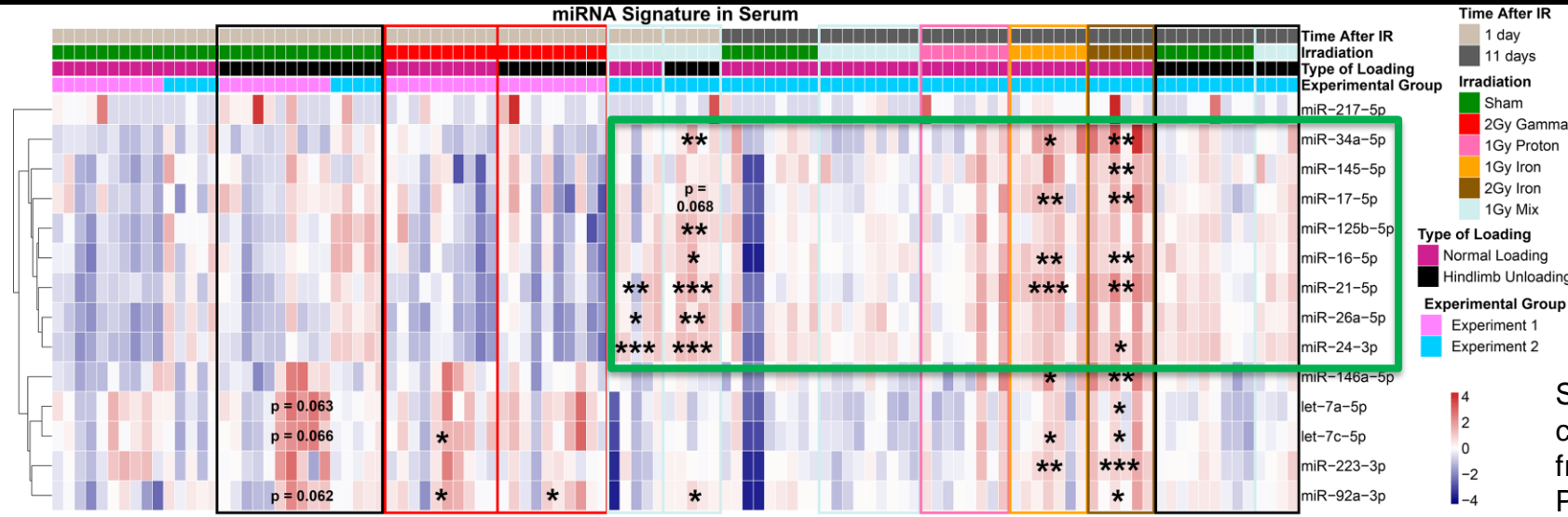
**Through ddPCR we are able to get exact counts of circulating miRNA in the serum**



Every single blue point on the plots represents one copy of miRNA (see arrows).

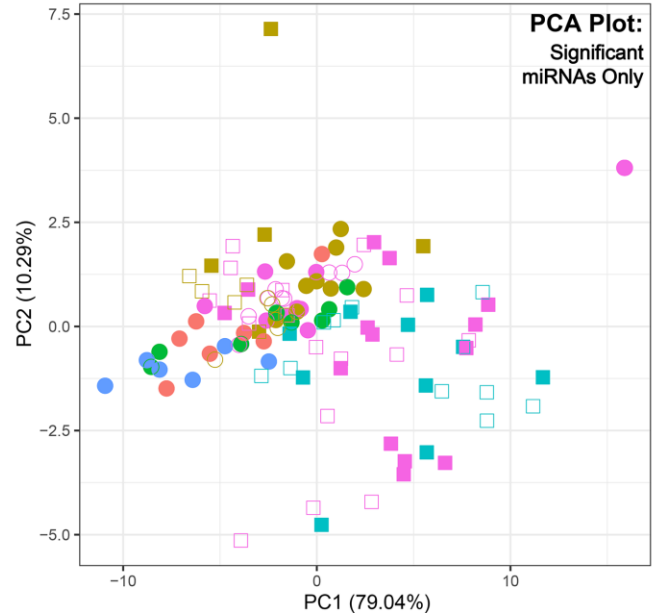
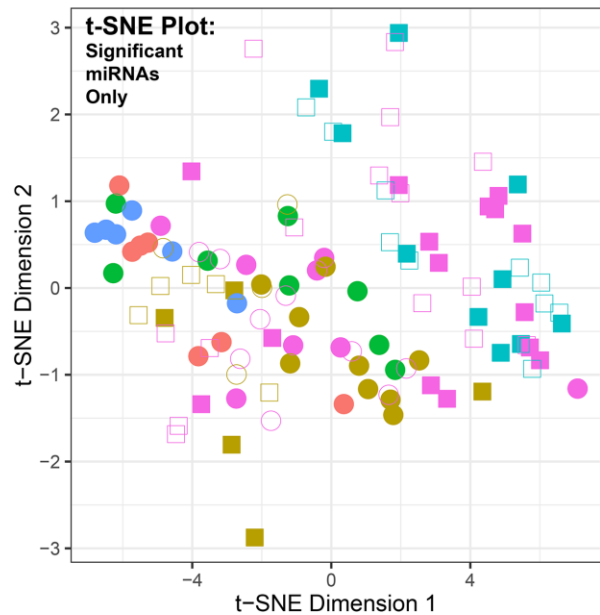
# Presence of miRNA signature in Serum of Mice in Simulated Space Environment

- HU for an initial three days followed by IR and continuation of HU for another 1 or 11 days
- Radiation exposure: Total body irradiation on conscious mice
  - 2Gy Gamma
  - 600 MeV/n <sup>56</sup>Fe (1 Gy and 2 Gy)
  - 150 MeV Proton (1Gy)
  - '1Gy Mix' (0.5Gy <sup>56</sup>Fe and 0.5Gy Proton)

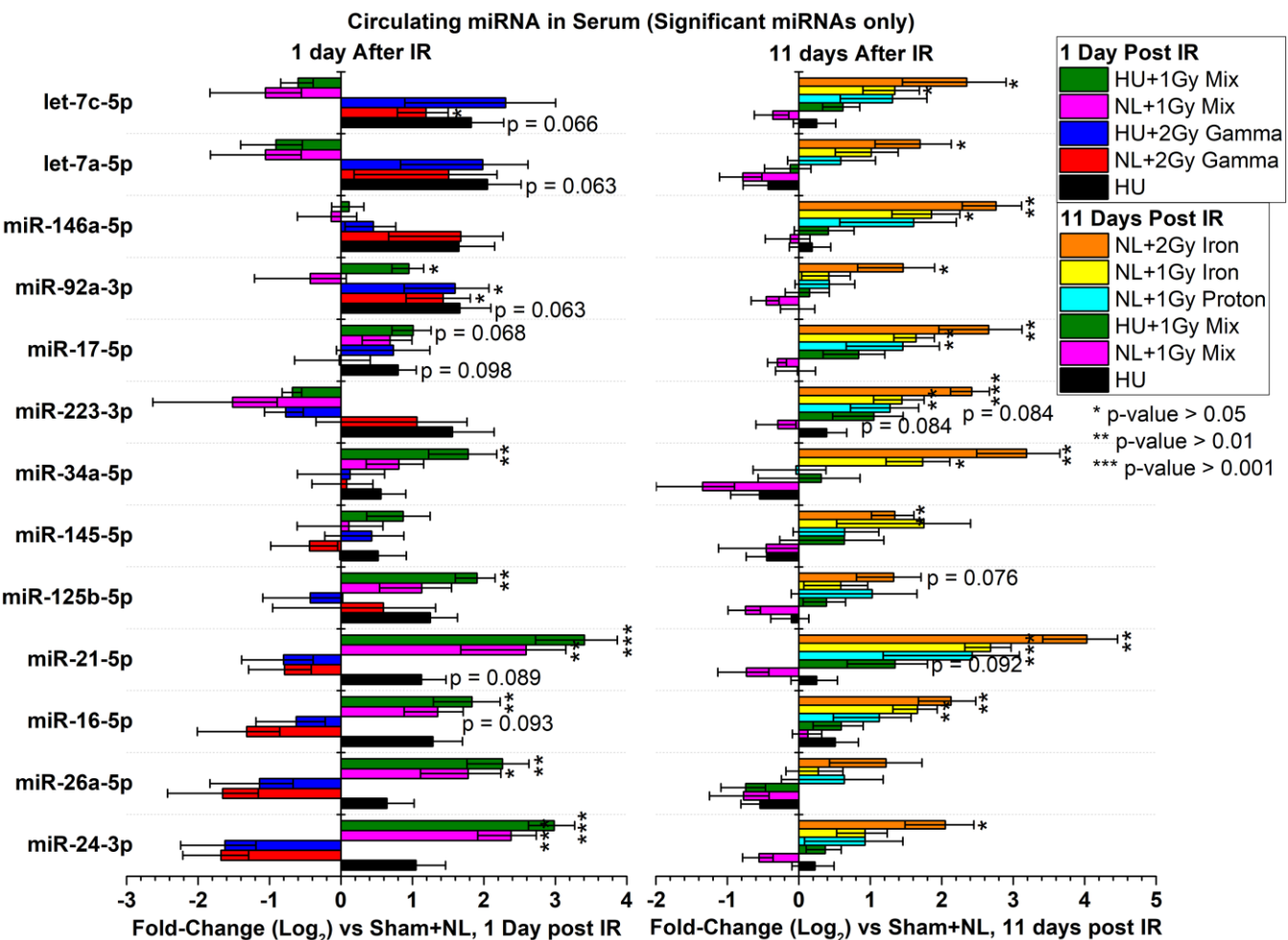


p-values determined by comparing to Sham Normal Loading Conditions, 1 day after IR

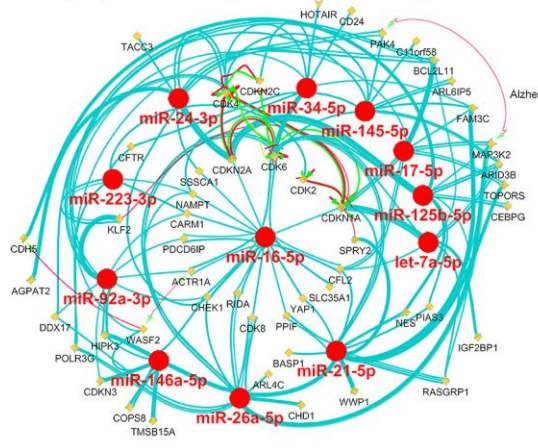
p-values determined by comparing to Sham Normal Loading Conditions, 11 days after IR



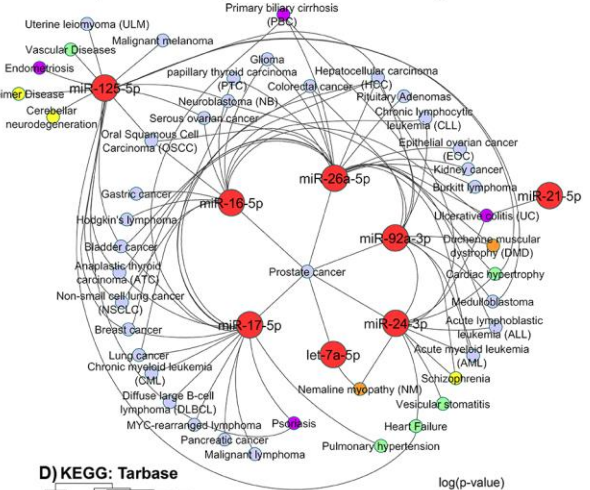
- Type of Loading: ○ HU, ● NL
- Irradiation: ● Sham, ● 2Gy Gamma, ● 1Gy Proton, ● 1Gy Iron, ● 2Gy Iron, ● 1Gy Mix
- Time Post IR: ● 11days, ■ 1day



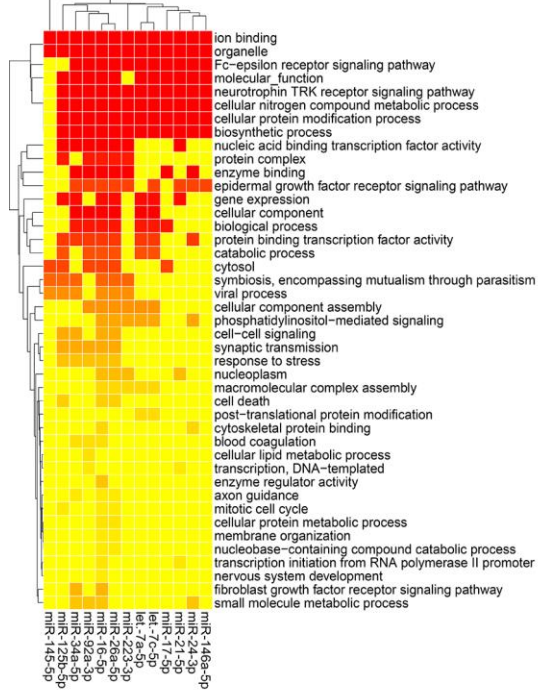
A) Top 50 Gene Targets for the miRNA Signature



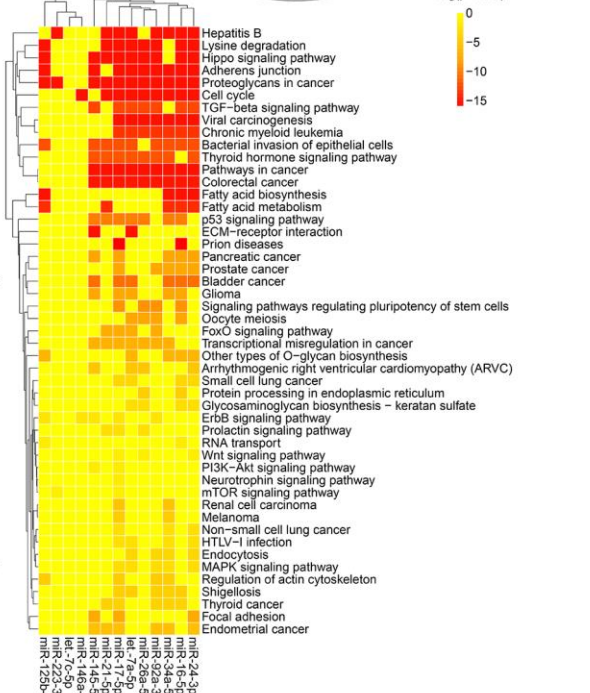
B) Diseases Regulated by the miRNA Signature



C) GO: microT-CDS



D) KEGG: Tarbase



**Confirmation exists in the miRNAs from the NASA Twin Study!!!**

## HUMAN EXPLORATION

NASA's Path to Mars

**EARTH RELIANT**  
MISSION: 6 TO 12 MONTHS  
RETURN TO EARTH: HOURS

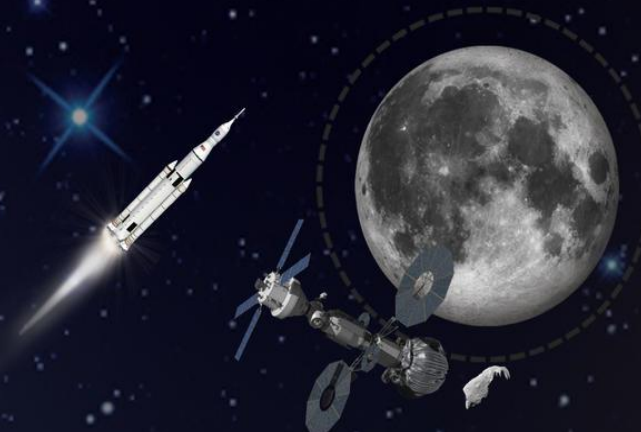
**PROVING GROUND**  
MISSION: 1 TO 12 MONTHS  
RETURN TO EARTH: DAYS

**MARS READY**  
MISSION: 2 TO 3 YEARS  
RETURN TO EARTH: MONTHS



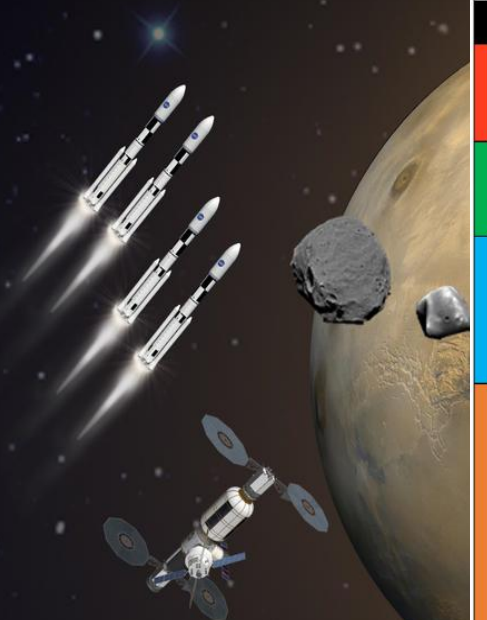
Mastering fundamentals aboard the International Space Station

U.S. companies provide access to low-Earth orbit

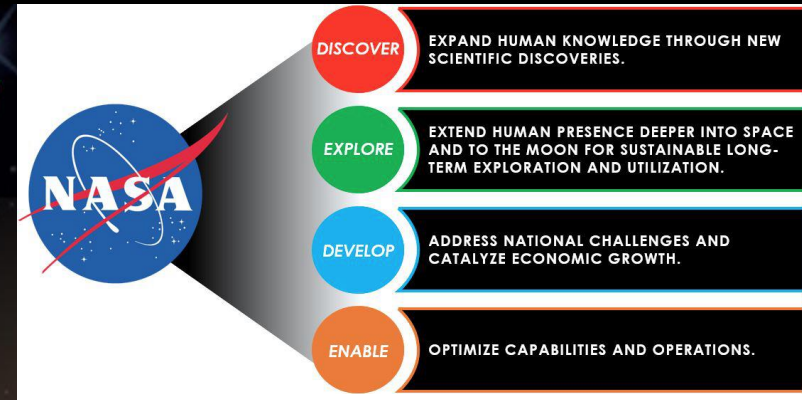


Expanding capabilities by visiting an asteroid redirected to a lunar distant retrograde orbit

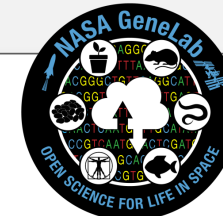
The next step: traveling beyond low-Earth orbit with the Space Launch System rocket and Orion spacecraft



Developing planetary independence by exploring Mars, its moons and other deep space destinations



NASA 2018 Strategic Plan Framework		
Theme	Strategic Goal	Strategic Objective
DISCOVER	EXPAND HUMAN KNOWLEDGE THROUGH NEW SCIENTIFIC DISCOVERIES.	1.1: Understand the Sun, Earth, Solar System, and Universe. 1.2: Understand Responses of Physical and Biological Systems to Spaceflight.
EXPLORE	EXTEND HUMAN PRESENCE DEEPER INTO SPACE AND TO THE MOON FOR SUSTAINABLE LONG-TERM EXPLORATION AND UTILIZATION.	2.1: Lay the Foundation for America to Maintain a Constant Human Presence in Low Earth Orbit Enabled by a Commercial Market. 2.2: Conduct Exploration in Deep Space, Including to the Surface of the Moon.
DEVELOP	ADDRESS NATIONAL CHALLENGES AND CATALYZE ECONOMIC GROWTH.	3.1: Develop and Transfer Revolutionary Technologies to Enable Exploration Capabilities for NASA and the Nation. 3.2: Transform Aviation Through Revolutionary Technology Research, Development, and Transfer. 3.3: Inspire and Engage the Public in Aeronautics, Space, and Science.
ENABLE	OPTIMIZE CAPABILITIES AND OPERATIONS.	4.1: Engage in Partnership Strategies. 4.2: Enable Space Access and Services. 4.3: Assure Safety and Mission Success. 4.4: Manage Human Capital. 4.5: Ensure Enterprise Protection. 4.6: Sustain Infrastructure Capabilities and Operations.





Sylvain Costes  
Space Bio  
miRNAs



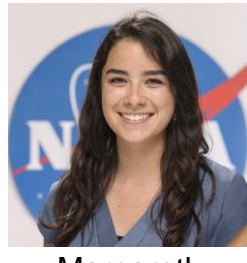
Egle Cekanaviciute  
Quantifying  
miRNAs



Sherina Malkani  
Quantifying miRNAs



Ann-Sofie  
Schreurs  
Provided Archived  
Tissues



Margareth  
Cheng-Campbell  
Quantifying  
miRNAs



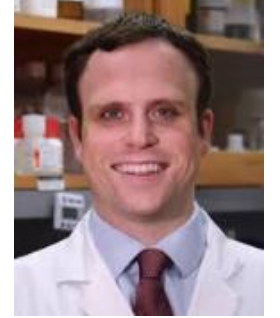
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Shirazi  
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Tissues



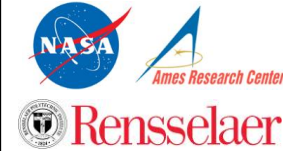
Ruth Globus  
Provided  
Archived  
Tissues



Elizabeth Blaber  
Space Bio miRNAs



Chris Mason  
Providing  
miRNA Twin  
Study Data



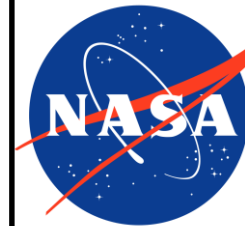
J. Tyson McDonald



Charles Vanderburg



Peter Grabham



This work is supported by:

16-ROSBFP GL-0005:  
NNH16ZTT001N-FG  
Appendix G: Solicitation of  
Proposals for Flight and  
Ground Space Biology  
Research

The Translational Research Institute  
through NASA Cooperative Agreement  
NNX16AO69A (T-0404)

