





The One-Year Mission

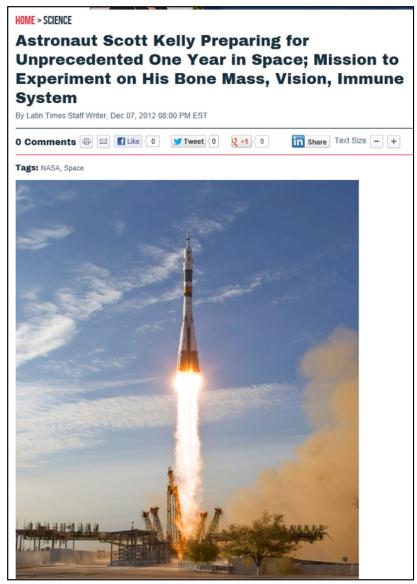


ISS Crew: Scott Kelly, Mikhail Kornienko Sign On For One-Year Mission



long space missions, such as voyages around the moon, to an asteroid and ultimately

to Mars, NASA officials said.



First Thoughts on A Twins Study





Great Value in n = 1 Longitudinal Omic Studies



Resource

Personal Omics Profiling Reveals Dynamic Molecular and Medical Phenotypes

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SIMMARY

Personalized medicine is expected to benefit from Personalized medicine aims to assess medical risks, monitor, combining genomic information with regular monitoring of physiological states by multiple highthroughout methods. Here, we present an integrative personal omics profile (IPOP), an analysis that combines genomic, transcriptomic, proteomic, metabolomic, and autoantibody profiles from a single individual over a 14 month period. Our iPOP analysis revealed various medical risks, including type 2 diabetes. It also uncovered extensive, dynamic changes in diverse molecular components and biological pathways across healthy and diseased contributors that can be difficult to assess. As such, the comconditions. Extremely high-coverage genomic bination of genomic information along with a dataled molecular and transcriptomic data, which provide the basis of our iPOP, revealed extensive heternallelic changes during healthy and diseased states and an unexpected RNA editing mechanism. This study demonstrates that longitudinal iPOP can be used to interpret healthy and diseased states by connecting genomic information with additional dynamic molecular constituents. For example, DNA microarrays have omics activity.

INTRODUCTION

diagnose and treat patients according to their specific genetic composition and molecular phenotype. The advent of genome sequencing and the analysis of physiological states has proven to be powerful (Cancer Genome Atias Research Network, 2011). However, its implementation for the analysis of otherwise healthy individuals for estimation of disease risk and medical interpretation is less clear. Much of the genome is difficult to interpret and many complex diseases, such as diabetes, neurological disorders and cancer, likely involve a large number of different genes and biological pathways (Ashiev et al., 2010; Grayson et al., 2011; Li et al., 2011), as well as environmental analysis of samples will be important for predicting, diagnosing and treating diseases as well as for understanding the onset, progression, and prevalence of disease states (Snyder et al., 2009).

Presently, healthy and diseased states are typically followed using a limited number of assays that analyze a small number of markers of distinct types. With the advancement of many new technologies, it is now possible to analyze upward of 10th allowed the subcategorization of Amphomes and gligmas

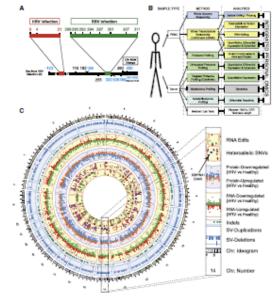
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Cell



Cell





(A)T ime countesummary. The subjectives monitored for a total of 726 days, during which there were two infections jied bar, HRV; geen bar, PGV). The black ba indicates the period when the subject (1) increased exercise, (5) inperied (1) mg of acetylasticytic acid and ibuprofen tablets each day (the latter only during the first 6 weeks of this period, and (i) substantially reduced sugar intake. Blue numbers indicate tasted time points

(B POP experimental design indicating the tissues and are uses involved in this study.

(Q Circos (Caywinski et al., 2009) pict summerbing (POP. From outer to inner rings: chamasome ideagram; genomic data (pale blue ring), structural vertants > 50 big id digitions (blue tiles), duplications in differs), index inner triangles; transpription is data (velow fro.), expression ratio of HFV infection to healthy states: proteomic data fight purple fing), ratio of protein levels during HRV infection to healthy status; transcriptomic data (yellow ring), differential heteroal expression ratio of alternative aliele to reference aliele for misserum and synonymous valants (burple dobs) and candidate RNA misserum and synonymous edits (md triangles, purple data, arange triangles and green data, respectively). See also Floure St.

WGS-Based Disease Risk Evaluation

We identified variants likely to be associated with increased susceptibility to disease (Dewey et al., 2011). The lat of high alleles are summarized in Figure 24), and 11 were validated by confidence SNVs and indeb was analyzed for rare alleles (<5% of the major silele frequency in Europeans) and for changes in genes with known Mendelian disease phenotypes (data summarized in Table 2, revealing that 51 and 4 of the rare coding SNV aplastic anemia (Yamaguchi et al., 2005), and (3) variants assoand indeb, respectively, in genes present in OMIM are predicted

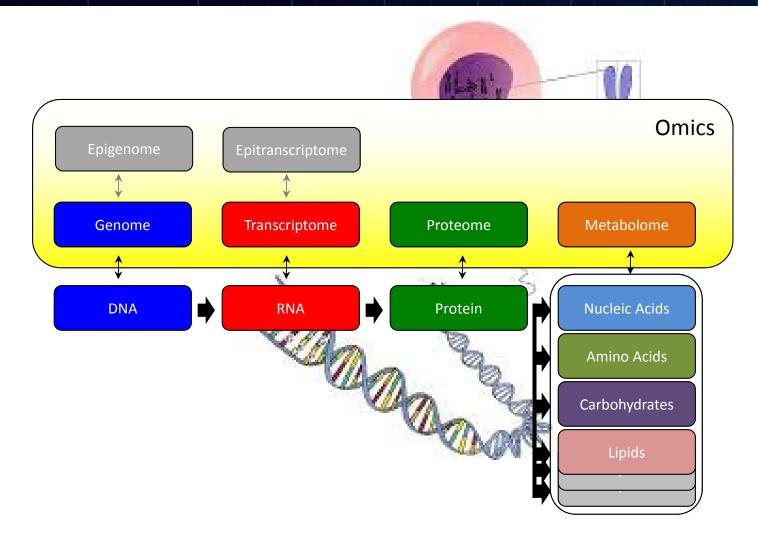
to lead to loss-of-function (Table S2A). This list of genes was further examined for medical relevance (Table S2A; example Sanger sequencing. High interest genes include: (1) a mutation (E366K) in the SERPINA1 gene previously known in the subject, (2) a damaging mutation in TERT, associated with acquired cisted with hypertriglyceridemia and diabetes, such as GCKR

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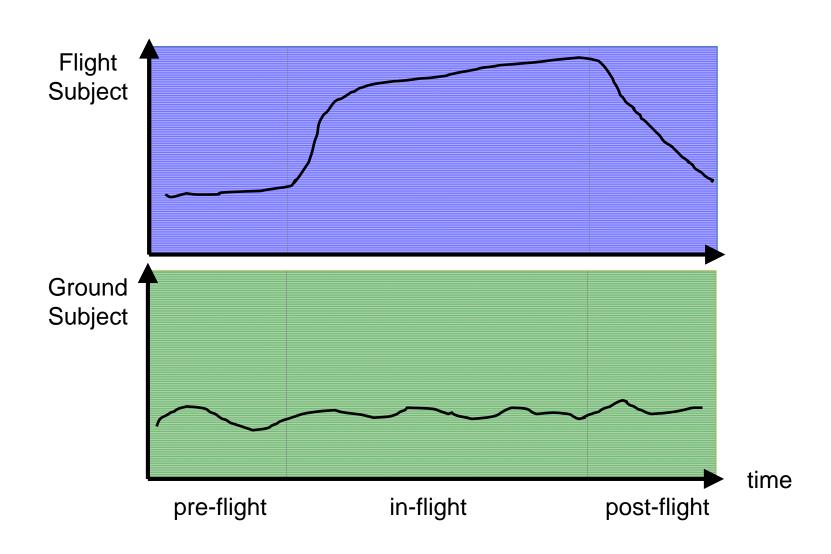
What is Omics?





One Notion





NRA Solicitation





National Aeronautics and Space Administration Johnson Space Center Human Exploration and Operations Mission Directorate Human Research Program Houston, TX 77058

Human Exploration Research Opportunities (HERO)

Appendix D

Differential Effects on Homozygous Twin Astronauts Associated with Differences in Exposure to Spaceflight Factors

Response Period: July 30, 2013 – September 17, 2013 Proposals Due: September 17, 2013, 5 PM Eastern Time Estimated Selection Announcement: January 2014

Appendix D - 1

"To capitalize on this unique opportunity,

NASA's Human Research Program (HRP) and the National Space Biomedical Research Institute (NSBRI) are initiating

a pilot demonstration project focused on the use of integrated human -omic analyses to

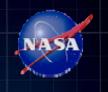
better understand the biomolecular responses to

the physical, physiological, and environmental stressors

associated with spaceflight."

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Selections



9

- 2 Subjects
 - Scott Kelly
 - Mark Kelly
- 10 Selections
 - 1. Susan Bailey, Colorado State University, Differential effects on telomeres and telomerase in twin astronauts associated with spaceflight
 - 2. Andrew Feinberg, Johns Hopkins University School of Medicine, Comprehensive whole genome analysis of differential epigenetic effects of space travel on monozygotic twins
 - 3. Christopher Mason, Weill Medical College of Cornell University, The Landscape of DNA and RNA Methylation Before, During, and After Human Space Travel
 - 4. Scott Smith, NASA Johnson Space Center, Biochemical Profile: Homozygous Twin control for a 12 month Space Flight Exposure
 - 5. Emmanuel Mignot, Stanford University School of Medicine, HERO Twin Astronaut Study Consortium (TASC): Immunome Changes in Space
 - 6. Stuart Lee, Wyle Laboratories, Metabolomic And Genomic Markers Of Atherosclerosis As Related To Oxidative Stress, Inflammation, And Vascular Function In Twin Astronauts
 - 7. Brinda Rana, University of California, Proteomic Assessment of Fluid Shifts and Association with Visual Impairment and Intracranial Pressure in Twin Astronauts
 - 8. Mathias Basner, University of Pennsylvania School of Medicine, HERO Twin Astronaut Study Consortium (TASC) Project: Cognition on Monozygotic Twin on Earth
 - 9. Fred Turek, Northwestern University, HERO Twin Astronaut Study Consortium (TASC) Project: Metagenomic Sequencing of the Bacteriome in GI Tract of Twin Astronauts
 - Michael Snyder, Stanford University, HERO Twin Astronaut Study Consortium (TASC) Project: Longitudinal integrated multi-omics analysis of the biomolecular effects of space travel



Differential effects on Telomeres and Telomerase in Twin astronauts associated with spaceflight





Susan Bailey, Ph.D. Colorado State Univ



Kerry George Wyle Labs/JSC

Specific Aims

The rate at which telomeres shorten provides an informative biomarker of aging and age-related pathologies (e.g., cardiovascular disease and cancer) that captures the interplay between genetics and lifestyle factors.

We propose that for the astronauts telomere maintenance is particularly relevant, as it reflects the combined exposures (e.g., radiation) and experiences (nutritional, psychological and physical stressors) encountered during space travel.

The Twins study provides the extraordinary opportunity to control variables of individual genetic differences, susceptibilities and lifestyle factors, making differential effects observed between the twins space-flight specific.

Comparisons with unrelated astronauts (separate study), will allow evaluating role of genetics/individual susceptibilities.

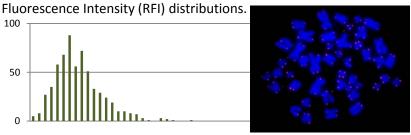
Our goal is to assess changes in telomere length and telomerase activity associated with the upcoming yearlong ISS mission in the space- and earth-bound twin astronauts.

We hypothesize that accelerated telomere shortening and elevated telomerase activity will be associated with space flight as compared to ground based control, in a duration and severity dependent manner.

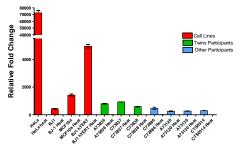
- Blood samples will be taken pre-flight (to establish baseline), in-flight (to evaluate short-term/temporary changes) and post-flight (to evaluate long-term/permanent changes)
- Data sharing for other endpoints will also inform this effort
- *In vitro* studies will investigate potential mechanisms (e.g., oxidative stress) and mitigation strategies (e.g., antioxidants)

Telomere length will be assessed using TELO-FISH

Florescence *in situ* Hybridization (FISH) with telomere probe on chromosomes (and interphase nuclei) is evaluated as Relative



Telomerase activity will be assessed using qRT-PCR TRAP quantitative Real Time-PCR Telomere Repeat Amplification Protocol



Implications of Research for Space & Earth



Space: This twins study will identify space-flight specific factors that influence telomere length and telomerase activity, informative biological indicators of aging and age-related degenerative diseases (e.g., cardiovascular disease and cancer). Our mechanistic investigations will begin to establish relevant relationships and suggest potential mitigation strategies for future study and to improve astronaut overall health.



Earth: Aging and age-related diseases like cardiovascular disease and cancer are an everyday concern on earth as well, therefore this study also seeks to make comparisons with unrelated astronauts (and controls) that will serve to identify individual susceptibility factors that influence telomere length and telomerase activity. Taken together with our mechanistic studies, mitigation strategies will be improved and applicable to all.

Epigenomics

NASA



Andrew Feinberg, M.D., M.P.H., and Jason Feinberg

Specific Aims

Aim 1. We will measure DNA methylation and chromatin at a genome-wide level in biological samples obtained from the space traveler at quarterly intervals, pre- and post-flight, and at times of unexpected exposures such as radiation events, or spacecraft environmental contamination. We also obtain measurements of the ground-based twin.

Aim 2. We will integrate epigenomic data with exposure to spaceflight conditions, looking for exposure-linked changes, and by comparison to the ground-based twin, determine whether these are transient or long-lived effects. We will also determine whether DNA mutations arise secondarily to these epigenetic changes.

Sample Collection and Analysis FD45 Fb190 FD223 FD350 FD0 Ground or ASAP RV and Ground Ground around ASAP. Blood mononuclear cells, buccal wash, at all time points Whole genome DNA sequencing prior to launch and post-recovery Whole genome bisulfite sequencing at several time points, 450K between

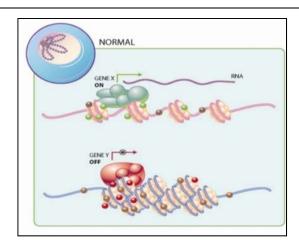
Implications of the Research for Space & Earth



Space: Identify reversible causes of genomic damage in space, e.g. radiation or toxin induced epigenomics change; quantify aging and genomic exposure.



Earth: First human study of the epigenome over time in a defined/controlled environment.



DNA methylation

ChIP-seg at all time points

RNAseq at several time points, arrays between

- Histone modifications (>200 known)
- Chromatin factor complexes
- Chromatin structure

Landscape of DNA and RNA Methylation



Christopher Mason, Ph.D.



Francine
GarrettBakelman,
M.D. Ph.D.

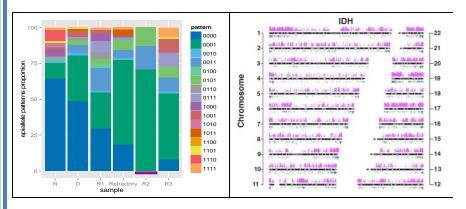
Epigenome Epitranscriptome Epiproteome

DNA NA rotein

#1 – Genome-wide epigenetic profiles of DNA methylation changes #2 – A comprehensive catalog of coding and noncoding, small and large RNA

#3 – Transcriptome-wide maps of RNA methylation sites

∆ in Epigenetics : Loci, regions, and clones



<u>A in Transcriptome</u>: Genes, Isoform, Edits, Allele, SNVs, ncRNAs, Fusions, & Methylation

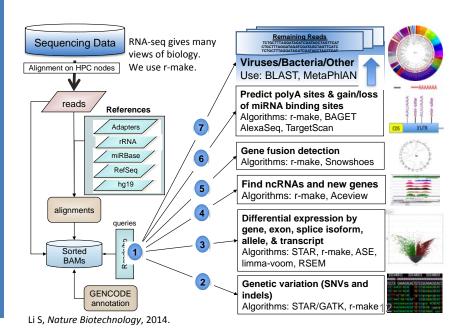
Implications of the Research for Space & Earth



Space: (1) Establish the genetic networks and expression patterns activated by space travel, (2) trace clonality of epigenetic changes, (3) examine the methylation of RNA



Earth: Aid research on aging, cancer, RNA biology, and circadian rhythm, all of which show differences at the (epi)genome & (epi)transcritpome



13 Jai

Biochemical Profile:

Homozygous Twin control for a 12 month Space Flight Exposure





Scott M. Smith, Ph.D.

Specific Aims

To provide a database of biochemical analyses from blood and urine samples. The analyses reflect a broad set of nutritional and physiological variables that may be altered as a result of the space flight environment (including diet, stress, weightlessness). Collecting data on the Ground twin will allow for a more direct comparison of the effects of space flight on human biochemistry and physiology.

Blood and urine collections **Preflight**:

L-180, L-45, L-10

In-flight:

FD15, 30, 60, 120, 180, 240, 300,

360

Post flight:

R+0, R+30





Space:

Improve understanding and time course of biochemical changes during flight and how the changes relate to diet during flight.



Earth:

Improve understanding of how diet can impact different biological systems.



Immunome studies in space





Emmanuel Mignot, MD, PhD

Specific Aims

- Study how long term space travel affects the immune system
- We will study how parameters of the immune system change at baseline and after a seasonal flu vaccination
- To do so, we study baseline and post flu parameters before, during and after a one year space flight



Stanford University

Implications of the Research for Space & Earth



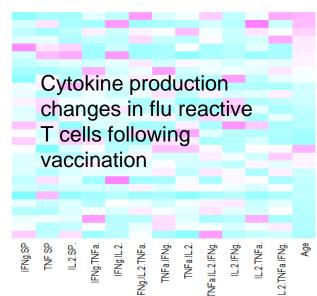
Space:

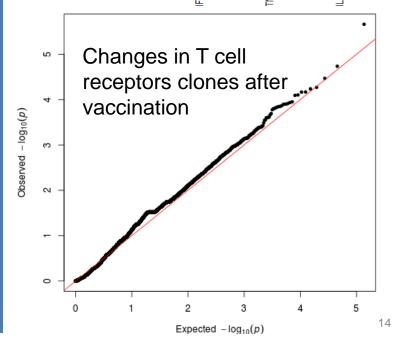
Will ensure that astronauts keep a healthy immune system during long flight, and stay protected against infections from earth when visitors are coming.



Earth:

Understand how immune response to vaccination differ in twins





METABOLOMIC AND GENOMIC MARKERS OF ATHEROSCLEROSIS IN TWIN

ASTRONAUTS

Specific Aims

- To study the effects of longduration spaceflight on the cardiovascular system independent of genotype
- To investigate relationships between gene expression, metabolomic profiles, biomarkers in blood and urine, and arterial structure and function using the space-flown and the ground-based identical twin

Stuart Lee

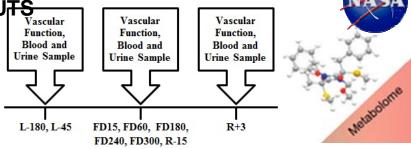
Implications of the Research for Space & Earth



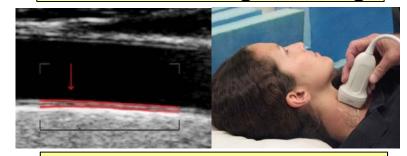
Space: Determine if the spaceflight environment perturbs genomic and metabolomic profiles and accelerates development of atherosclerosis (occupational health)



Earth: Develop novel insights of how longitudinal changes in genomic and metabolomic profiles are related to risk factors for atherosclerosis



Pre- and Post-flight Testing



Inflight Operations





PROTEOMIC ASSESSMENT OF FLUID SHIFTS AND ASSOCIATION WITH VISUAL IMPAIRMENTS AND INTRACRANIAL PRESSURE IN TWIN ASTRONAUTS





Brinda Rana, PhD Vivian Hook, PhD

Specific Aims

To explore proteomic and genomic biomarkers underlying space flightinduced fluid shifts and visual Mike Stenger, Pho impairment & intracranial pressure (VIIP) symptoms.



The proteome is the set of proteins produced by the genome at a given time. Proteomics captures the state of molecular and cellular processes at a specific time point.

Implications of the Research for Space & Earth



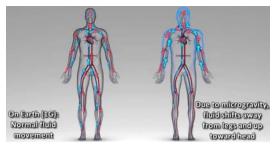
Space: Discovery of molecular pathways involved in the evolution of spaceflight adaptations related to fluid redistribution inflight and the etiology of visual acuity and ocular changes in-fight and post-flight.



Earth: This project has broader impact on Earth-based clinical areas such as traumatic brain injury-induced elevations of intracranial pressure, hydrocephalus, and glaucoma







In-flight Operations



Blood Plasma collection Ultrasound measures of fluid shifts Intracranial Pressure Intraocular Pressure Ocular Structure **Blood Pressure Heart Rate** Vascular Resistance

Pre- and Post-flight Testing



All in-flight operations and: Tissue hydration **MRI**

16

Cognitive Performance in Spaceflight

Specific Aims

There are a number of environmental stressors unique





Mathias Basner M.D., Ph.D.

to the spaceflight environment that may affect cognitive performance, which is crucial for mission success. Our main objective in the TWINS study is to investigate whether cognitive performance is affected by initial and prolonged exposure to the spaceflight environemnt and after return to Earth. We will use the Cognition test battery, which consists of 10 brief neuropsychological tests that were specifically designed for high performing astronauts. We will compare data within subjects, between twins, relative to astronauts flying 6-month missions, and relative to normative data gathered in astronauts on the ground. The cognitive data will be correlated with markers derived from biological samples taken before, during, and after the 12-month mission.



Ruben C. Gur. Ph.D.

Implications of the Research for Space & Earth



Space ploration-type missions require humans to spend unprecedented durations in space, yet our knowledge on the effects of prolonged exposure to the spaceflight environment is very limited. After the study, we will have an initial understanding of whether and to what extend prolonged ISS missions are associated with changes in cognitive performance, and how these relate to biologic markers.



Earth The results have direct implication for other high performing populations exposed to stressful environments for prolonged periods of time on Earth.

	Test	Test Cognitive Domain Brain Regions (from fMRI studies)				
фV	Motor Praxis (MPT)	Sensory-motor ability	Sensorimotor Cortex	0.51		
	Visual Object Learning (VOLT)	Visual object learning and memory	Medial Temporal Cortex - Hippocampus	1.69		
	Fractal 2-Back (F2B)	Attention and working memory	Dorsolateral prefrontal Cortex, Cingulate, Hippocampus	1.93		
	Abstract Matching Task (AMT)	Abstraction and mental flexibility	Prefrontal Cortex	2.33		
1	Line Orientation (LOT)	Spatial orientation	Right Temporo-Parietal Cortex, Visual Cortex	2.07		
(a)	Emotion Recognition (ERT)	Emotion recognition	Cingulate Cortex, Amygdala, Hippocampus, Fusiform Face Area	2.03		
?	Matrix Reasoning (MRT)	Abstract reasoning	Prefrontal Cortex, Parietal Cortex, Temporal Cortex	2.09		
Т	Digit Symbol Substitution (DSST)	Complex scanning, visual tracking, attention	Temporal Cortex, Prefrontal Cortex, Motor Cortex	1.60		
	Balloon Analog Risk (BART)	Risk decision making	Orbital frontal Cortex, Amygdala, Hippocampus, Anterior Cingulate Cortex	2.39		
200	Psychomotor Vigilance (PVT)	Vigilant attention and psychomotor speed	Prefrontal Cortex, Motor Cortex, Visual Cortex	3.17		

The Cognition Test Battery

Cognition was specifically designed for astronauts and is currently used during 6-month ISS missions and in multiple space analog environments (including Antarctica, HI-SEAS, and HERA).

The Bacteriome in the Gastrointestinal Tract



Fred Turek, Ph.D.

Specific Aims

The GI tract of humans is populated by a diverse "ecosystem" of micro organisms, mostly bacteria: the bacteriome. The bacteriome can <u>help--</u>contributing to digestion and immune system function-- or <u>harm</u>-- overgrowth of some types accompanies illness or stress.

This project will examine what changes occur to the bacterial populations over a year in space, that are different from the changes over time on Earth. Are particular types of bacteria susceptible to the space environment, and if so, which types?

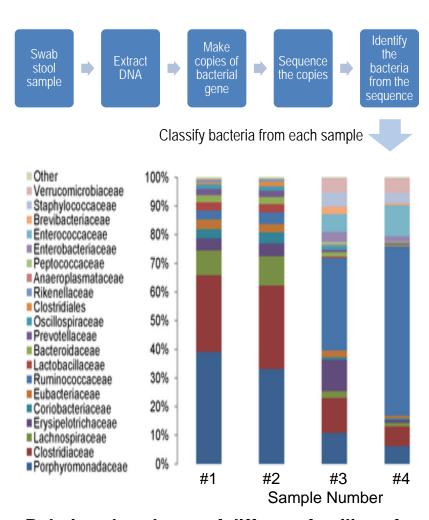
Implications of the Research for Space & Earth



Space: Knowing how the bacteriome changes over time in space can help us make plans to protect astronauts' health for longer-term space flights. For example, adjustments to diet could help maintain beneficial bacterial types.



Earth: Observing how the bacteriome changes in relation to health and environmental changes, (such as those studied in other Twin Projects) can provide insights into how the bacteriome may respond to challenges and contribute to the human host's health.



Relative abundance of different families of bacteria. Will there be systematic changes in the twin in space not seen in the twin on Earth?

Integrated Omics: Mike Snyder, Ph.D.





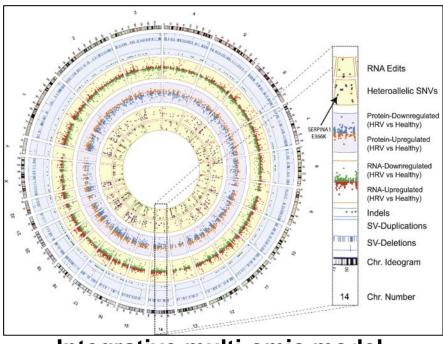
Michael Snyder, Ph.D



Juliane Winkelmann, M.D.

Specific Aims

Our main objective in the twin study is to perform a complete analysis of all biomedical and molecular data collected during the mission to produce the singular most comprehensive portrait of the human biophysical response to the rigors of spaceflight. We are at an unprecedented era in genomic medicine, allowing for the sensitive and precise measurement of billions of biochemical molecules, which will allow us to detect the subtlest of changes in Scott and Mark's physiology over time. By integrating these data, we can follow alterations in their cellular systems to both better understand the effects of space travel on human health, and how an astronaut's genome may contribute to his/her own unique physiologic response to microgravity.



Integrative multi-omic model

								1		_
						—	 		Dyslipidemia	1
					—				Coronary artery disease	30
			-						Basal cell carcinoma	6
		•		_					Type 2 diabetes	28
			-						Hypertriglyceridemia	4
			•						Osteoarthritis	1
•									Age related macular degeneration	12
	_	: 							Hypertension	5
	_	_							Obesity	11
	4								Asthma	13
	_								Stroke	4
_									Psoriasis	14
•			-							

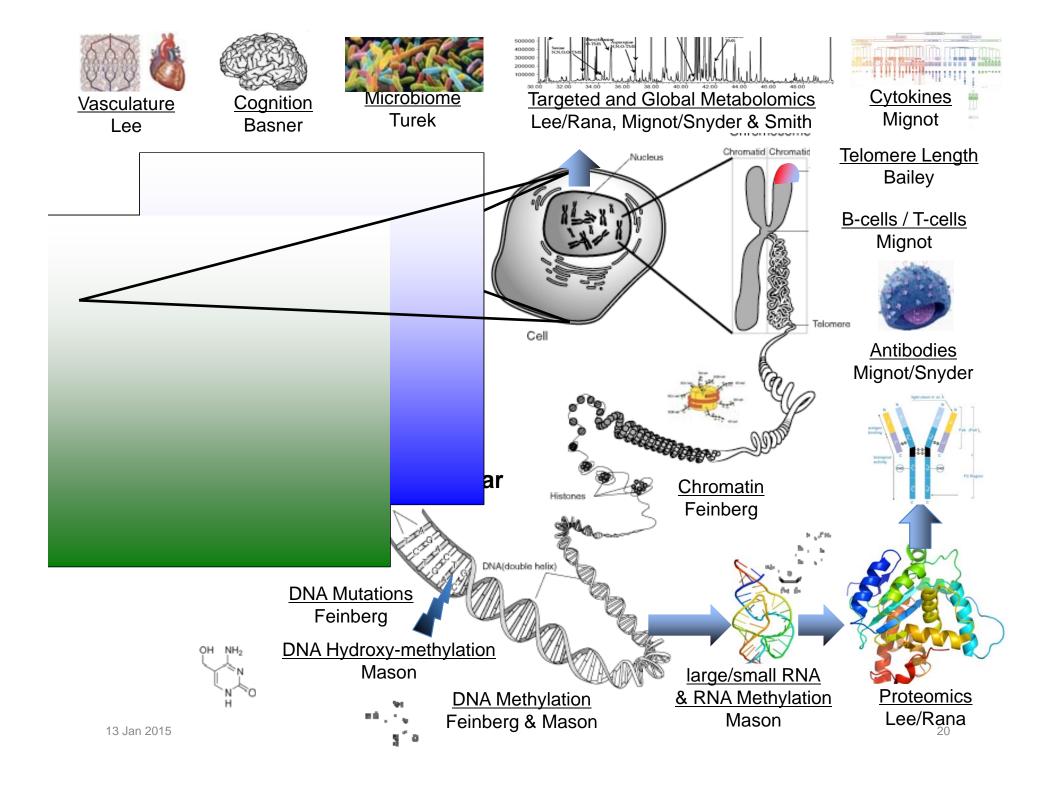
Implications of the Research for Space & Earth

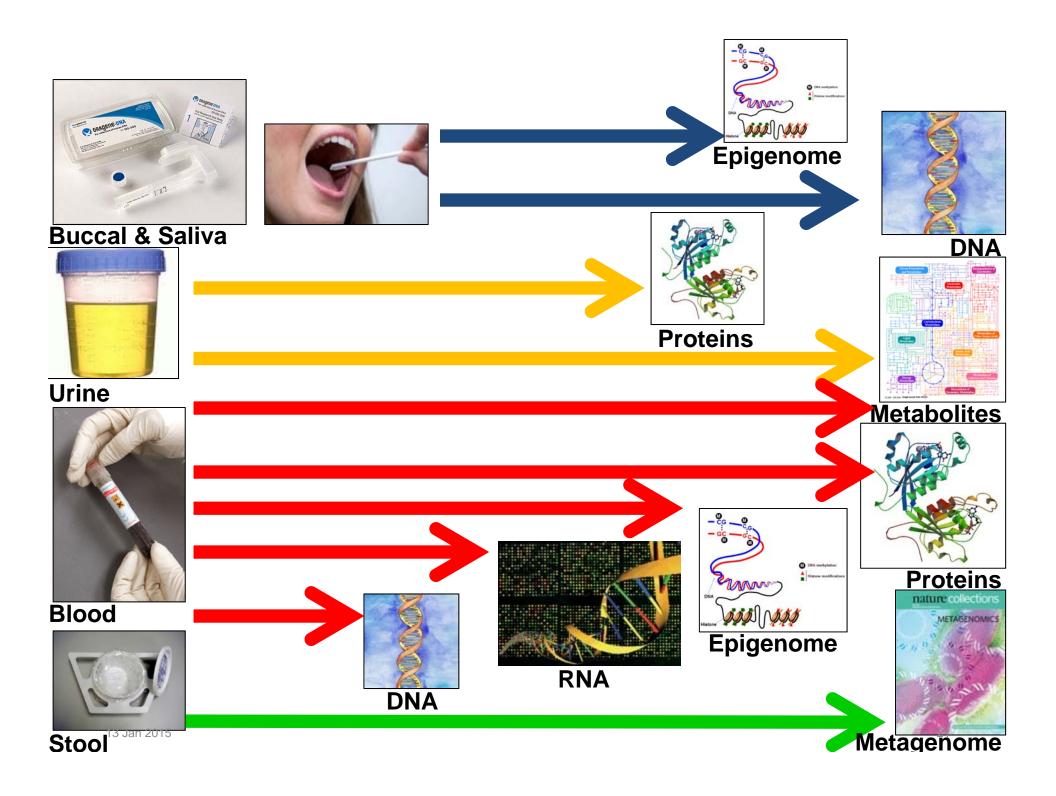


Space:We will generate a detailed benchmark for how human physiology changes in space in great molecular detail. This wealth of data will be essential for any future planning of long duration space exploration missions, and provide a proof-of-principle for better monitoring and managing astronaut health.



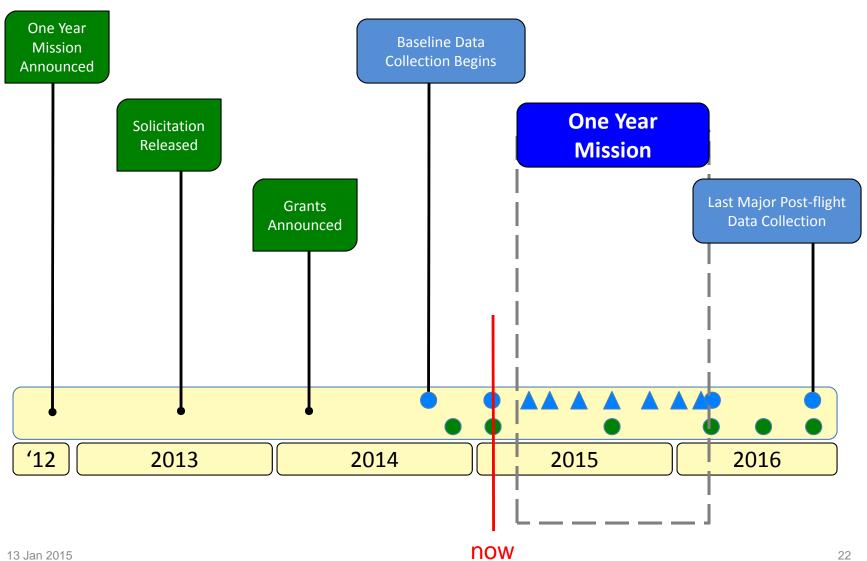
Earth:With this study, Scott and Mark Kelly will be the most thoroughly profiled twins in history, and the resultant data will offer new insights into how two siblings with nearly-identical genomes respond to different conditions.





Timeline





Issues Associated with Omic Research

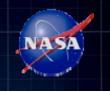


Research ethics

- The primary risks involved in genetic research are risks of social and psychological harm, rather than risks of physical injury
 - Could provoke anxiety and confusion about disease risk
 - Uncover unwanted information about heritage, ancestry, and family relationships
- De-identification of genomic information
- Information given to subjects
 - Individual genome sequence data?
 - Interpretation of the genome sequence and/or genetic counseling?
 - Option to decline to receive all or part of the results (Right Not to Know)?
- Researcher's access to genomic information
- Interim policy on genetic research JID 1800.4
- NASA policy anticipated summer 2015
- Medical care
- Occupational health
- Insurance (health, disability, life)
- Employment activity



Conclusion



 The Twins Study (Scott and Mark Kelly) is NASA's first foray into 21stcentury omics research

- Built around Scott Kelly's one year mission
- The Twins Study will examine
 - Genome, telomeres, epigenome
 - Transcriptome and epitranscriptome
 - Proteome
 - Metabolome
 - Physiology
 - Cognition
 - Microbiome
- NASA is addressing
 - Protections for research participants
 - Use of data in medical care, occupational medicine, mission planning



Acknowledgements





The Twins Study Investigator Team



John Charles



Mike Barratt



Bill Paloski



Graham Scott

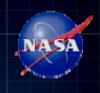


Jeff Sutton



Mark Shelhamer

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- Karen Lawrence
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- Pasha Morshedi
- Kelly Norwood
- Robert Pietrzyk
- Laura Sarmiento
- Clarence Sams
- Alissa Schade
- Nichole Schwanbeck
- Wesley Tarkington
- Jennifer Wilson



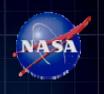


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