

MEETING ABSTRACT

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# Meta-analysis of gene expression changes in response to radiation exposure

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From 10<sup>th</sup> Annual UT-ORNL-KBRIN Bioinformatics Summit 2011  
Memphis, TN, USA. 1-3 April 2011

## Background

Given NASA's recent focus on long-duration space travel, potential adverse effects on astronauts of ionizing radiation need to be minimized. Levels of exposure for astronauts on such flights can be high enough to cause damage to DNA, possibly causing mutation and cancer. By analyzing the mRNA expression levels of genes exposed to different doses of gamma radiation in laboratory experiments, it can be determined which genes act as biomarkers of dosage-specific radiation exposure which can then be used on lab-on-a-chip (LOC) diagnostic platforms for early detection of ionizing radiation exposure.

## Methods

Currently, our group is incorporating three genes known to be involved in double-stranded DNA damage and repair, including: p21, p53, and  $\gamma$ H2AX. While these genes should show up in response to ionizing radiation exposure, they are not ideal biomarkers due to their lack of specificity. Therefore, by using meta-analysis, our goal is to gain further insight into additional biomarkers for radiation exposure. More than 40 publicly-available data sets appropriate to our study were obtained from the online repository, Gene Expression Omnibus (GEO) [1]. These gene expression experiments were subsequently categorized by radiation type and dose. Meta-analysis was performed using a combination of statistical tests using the Differential Expression via Distance Synthesis (DEDS) package [2].

## Results

Preliminary meta-analysis of these publicly available datasets yields potential biomarkers from the P53 signaling pathway (CDKN1A, GADD45A, MDM2, PMAIP1), stress response transcription factors (ATF3, JUN, JUNB, JUND), and cell surface receptors (CD69, CD70, CD83). Additional microarray experiments involving irradiated blood samples are underway. The analysis of both publicly available data and our own datasets will yield a broader picture of genes most sensitive to exposure of ionizing radiation for use as biomarkers on LOC diagnostic platforms for early detection of radiation exposure, leading to subsequent treatment.

## Acknowledgments

Members of the University of Louisville Research Group for Diagnosing and Mitigating Human Exposure to Radiation Using Micro-Nanotechnology include Robert W. Cohn, John Eaton, William D. Ehringer, Andre M. Gobin, Andrea S. Gobin, Balaji Panchapakesan, Eric C. Rouchka, Palaniappan Sethu, and Robert S. Keynton. This work was supported by the National Aeronautics and Space Administration (NNX10AJ36G) and the National Institutes of Health (NIH) (P2ORR016481, P30ES014443). Its content is solely the responsibility of the authors and does not necessarily represent the official views of NASA, NCCR, NIEHS, or NIH.

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Published: 5 August 2011

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doi:10.1186/1471-2105-12-S7-A20

**Cite this article as:** Kirtley *et al.*: Meta-analysis of gene expression changes in response to radiation exposure. *BMC Bioinformatics* 2011 **12** (Suppl 7):A20.

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