

(Phospho-)proteomic Signaling Responses of Human Male Germ Cell Lines to Simulated Microgravity and Hypogravity

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INTRODUCTION & AIM

Over the past century, advancements in space technology and the growing interest in space exploration, including the rise of space tourism, have significantly increased human participation to spaceflight. However, as human exploration extends beyond low Earth orbit (BLEO), with an increase in mission duration and complexity, there is a need for comprehensive research on the mechanisms of adaptation of human physiology to long-term exposure to altered gravitational conditions. Among the organ systems affected by microgravity, the reproductive one has been largely under investigated, despite evidence of disrupted endocrine signalling and altered gonadal cell function during spaceflight. Understanding the effects of microgravity on germ cells is crucial as any alterations could be inherited by offspring.

The present study aims to fill this gap by providing new insights into how altered gravity affects, at the molecular level, male reproductive health. To this end, we exposed two human testicular cancer cell lines to simulated microgravity (SμG) and hypogravity (ShG). Specifically, we selected the TCam-2 cell line, which resembles primordial germ cells/gonocytes, and the NT2D1 cell line, which exhibits pluripotency characteristics. A targeted (phospho-)proteomics approach, namely Reverse-Phase Protein microArrays (RPPAs), was exploited to investigate the molecular mechanisms that determine the cell response to an altered gravitational field.

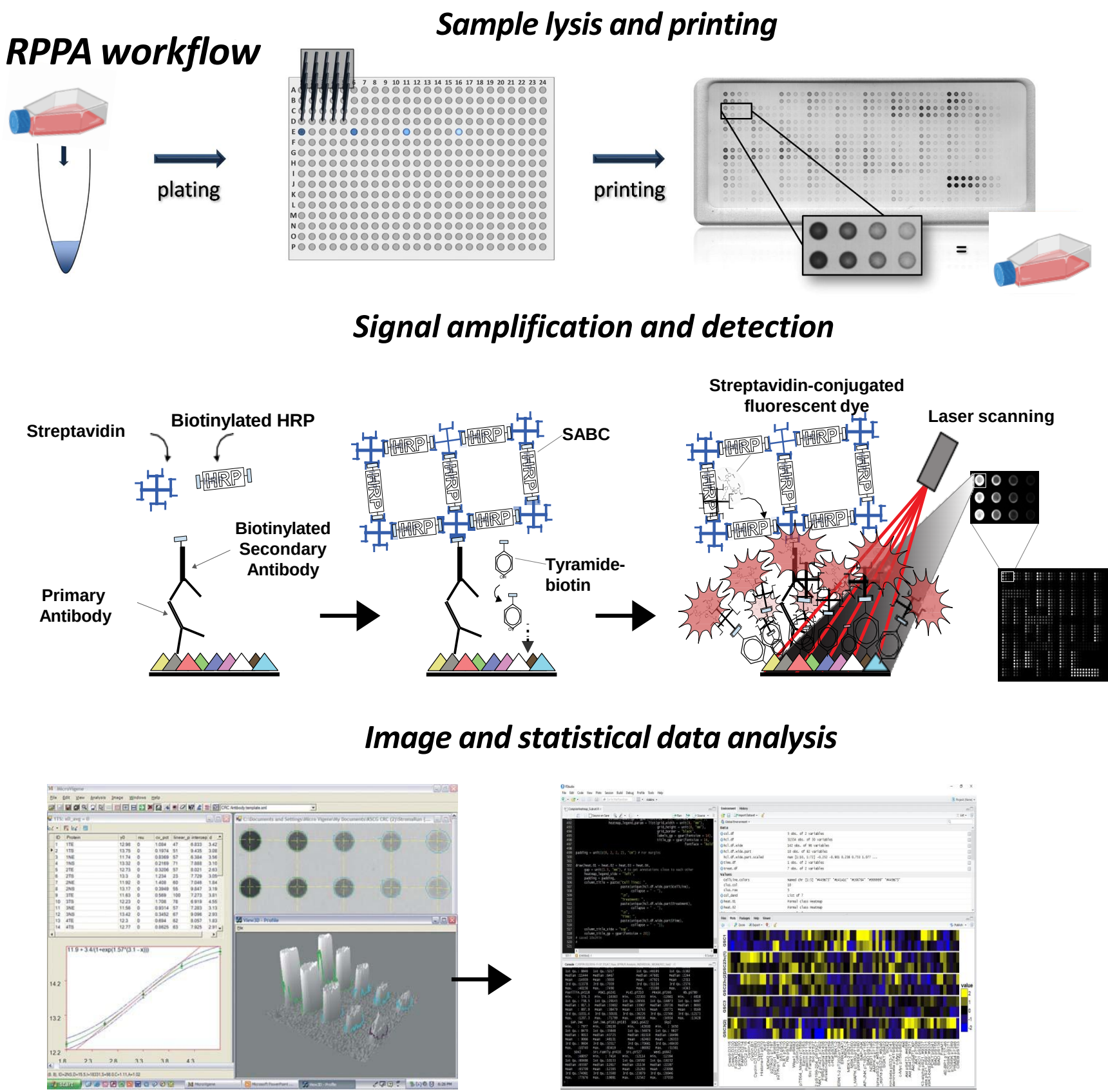
METHODS

Cell Culture

TCam-2 and NT2D1 cell lines were 2D cultured and exposed for 3, 24, and 72 hours to unitary gravity ("on ground": OG) or simulated microgravity (SμG) and hypogravity (ShG), using a Random Positioning Machine (RPM). According to the principle of "gravity-vector averaging" the RPM machine mimics the effects of SμG and ShG on standard cell culture flasks.

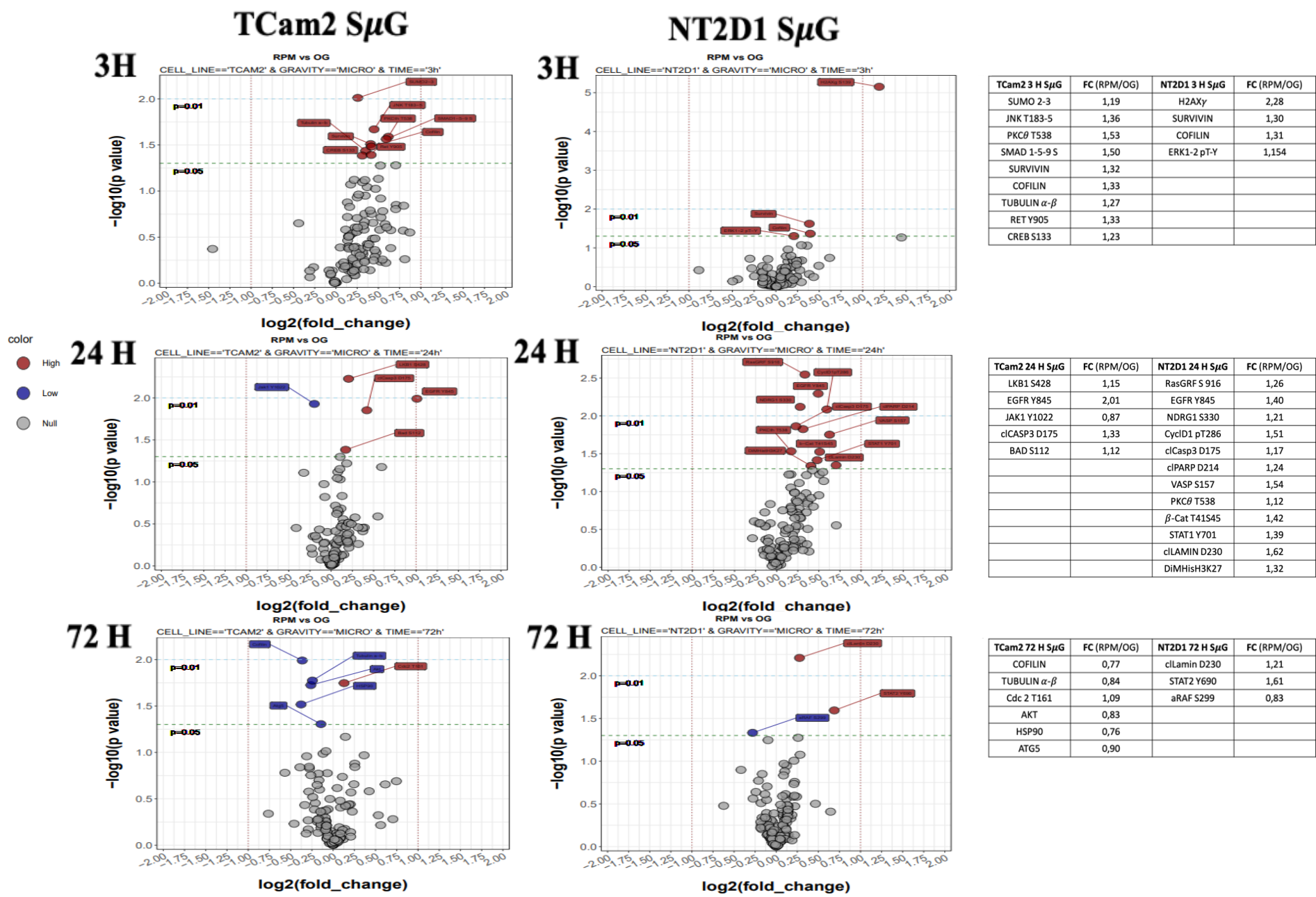
Reverse-Phase Protein microArrays (RPPA)

RPPA analysis was conducted using a panel of 130 antibody targets, potentially affected by altered gravity conditions, including but not limited to cytoskeleton, cell growth and proliferation as well as cancer-related pathways.

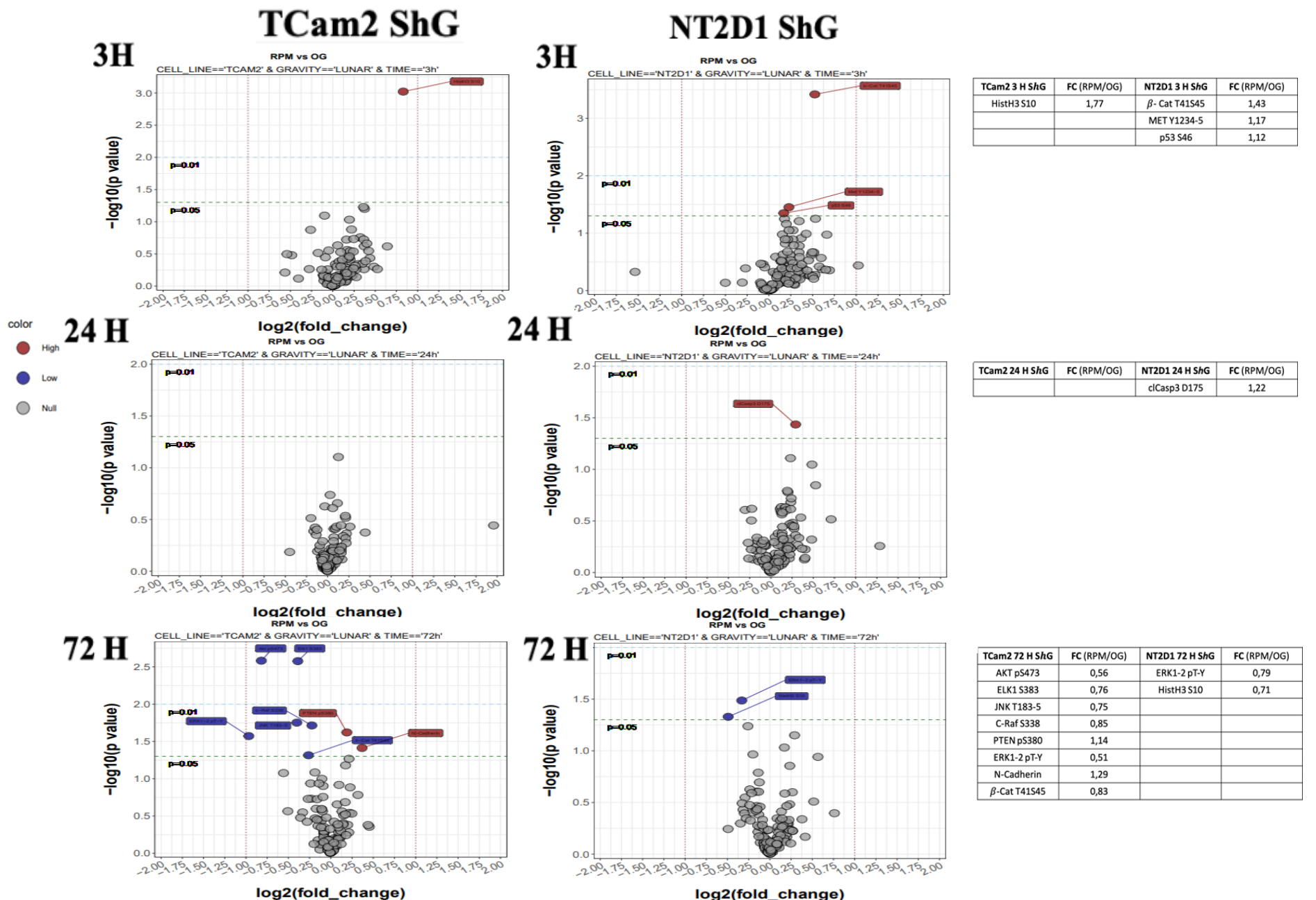


RESULTS & DISCUSSION

Panel 1. Exposure to SμG induces early upregulation of some phospho-proteomic pathways and late downregulation of some other phospho-proteins.



Panel 2. ShG has a limited impact on signaling pathways and mostly leads, in late exposure, to a decrease of some phospho-protein levels



The figure shows volcano plots of TCam-2 and NT2D1 exposed to SμG (panel 1) or ShG (panel 2) for the indicated timepoints. Dots represent the log2 of RPM to OG ratio (x axis) versus negative log10 of p value (y axis) for each individual antibody analyzed. Statistical significance was set to 0.05 (n=3). Statistically significant results (RPM versus OG) are labeled and colored as blue and red dots, respectively for down- and upregulated antibodies in the volcano plots and listed in a tabular format on the right of both Panel 1 and 2.

CONCLUSION

This study investigated the (phospho-)proteomic responses of TCam2 and NT2D1 germ cells to altered gravity conditions (simulated micro- and hypo-gravity) using RPPA. Overall, we found that modifications of the gravity vector result in a limited but significant number of phospho-protein changes (29/130 for TCam2 and 25/130 for NT2D1, respectively). These findings demonstrate the sensitivity of these cell lines to altered gravity confirming previous studies of our group on TCam-2 cells [Ferranti F. et al., Biomed Res Int. 2014; Berardini M. et al., Cells 2023]. Further investigations are needed to disclose whether such molecular changes are intrinsic features of the selected cancer cell lines or are a shared biological property of germ cells. Moreover, the transient nature of these molecular changes, and their difference over the exposure time, leaves an encouraging margin for cellular adaptation to altered gravitational force.

FUTURE WORK / REFERENCES

Confirmation of significant results via orthogonal techniques and complementation of RPPA data with other molecular analyses, in order to obtain a comprehensive scenario of the cellular responses to SμG and ShG.