

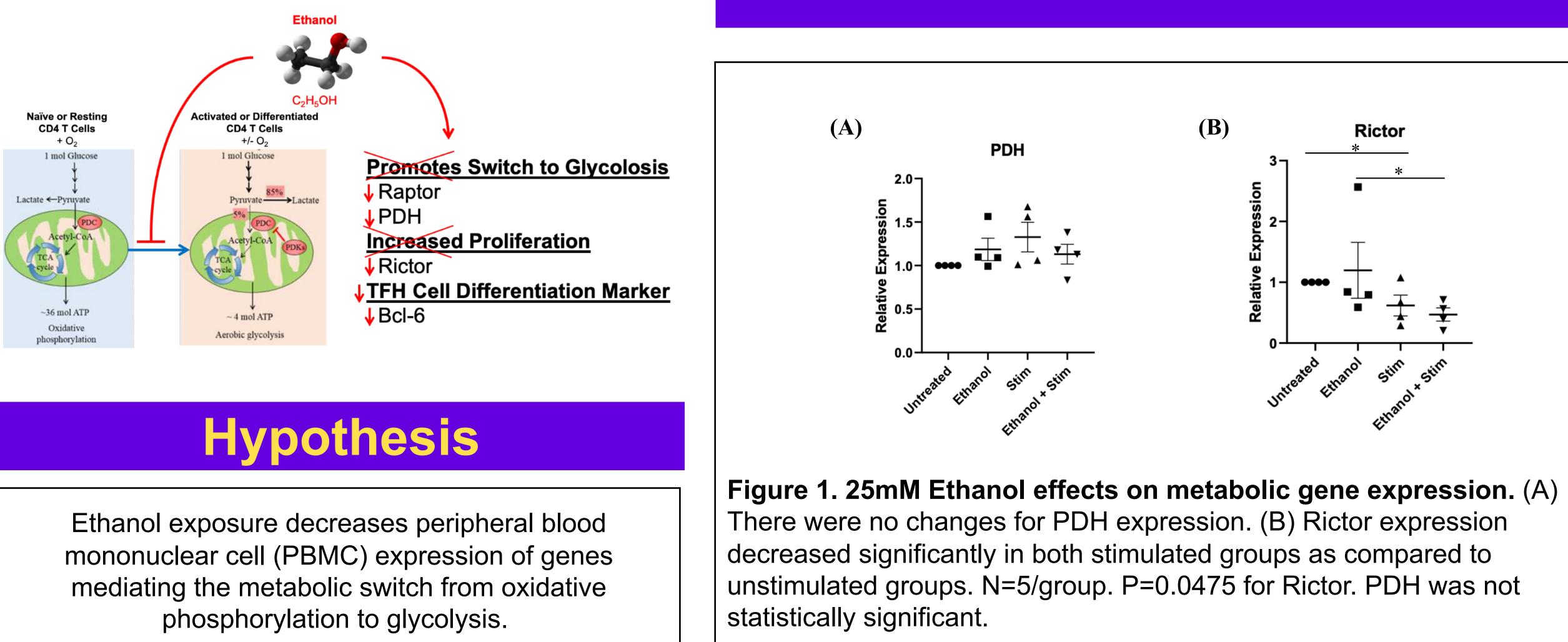
School of Medicine

Isabella Welsh, Patrick McTernan, Robert Siggins, Patricia E Molina Louisiana State University Health Science Center

Introduction

- Alcohol impairs the host defense immune response to infection by viruses, like HIV, and by a variety of bacteria.
- Published work has shown that acute exposure to ethanol decreases immune cell activation and proliferation both in preclinical animal models and *in vitro* cell culture.
- Recently, the importance of metabolic control of immune activation and differentiation has been described.
- Optimal proliferation and differentiation requires a metabolic switch from the efficient, but slow, production of ATP by oxidative phosphorylation to more rapid ATP generation by glycolysis (at the expense of efficiency).
- However, the mechanistic role of this metabolic switch in ethanol's effects on immune cell activation, proliferation, and differentiation is unknown.

Conceptual Model



Effect of Acute Alcohol on Peripheral Blood Mononuclear Cell Metabolism

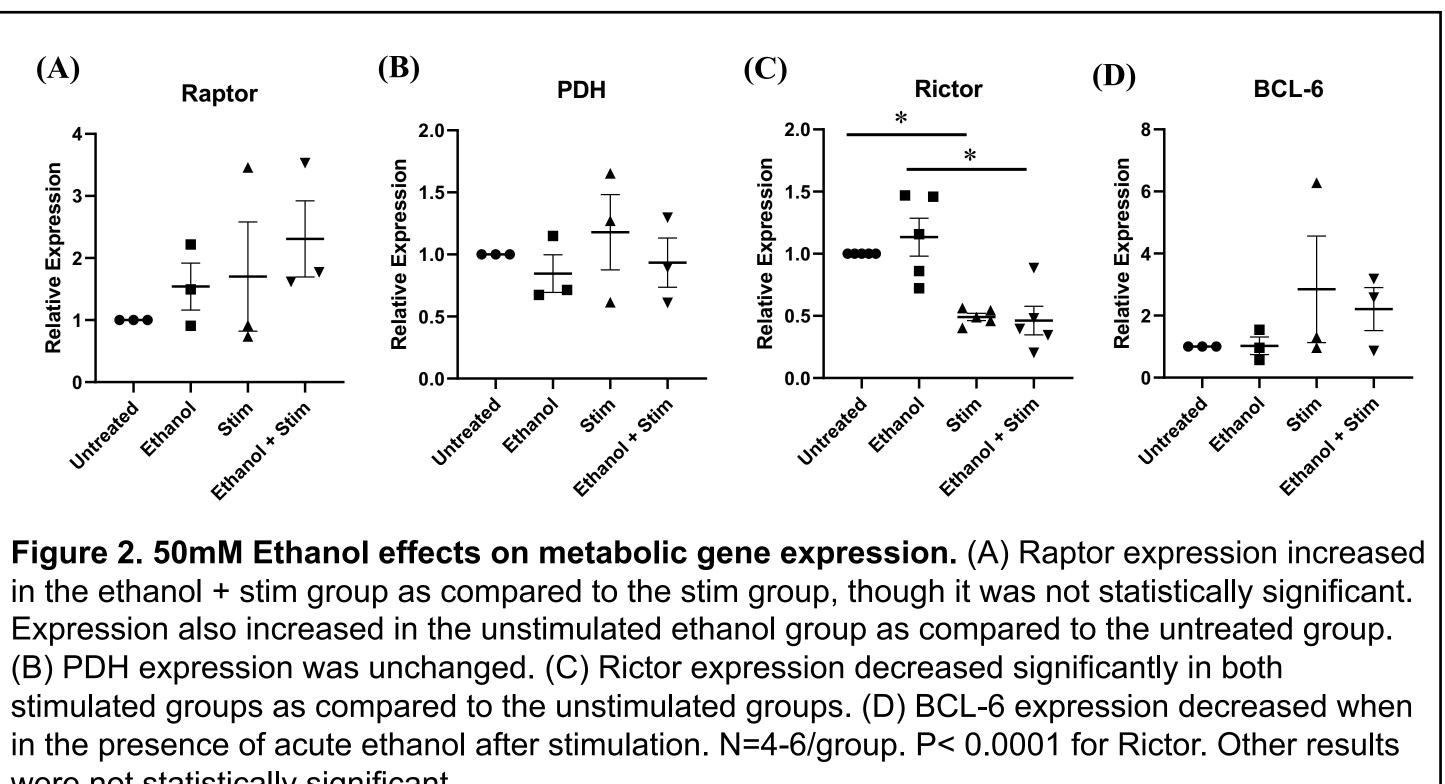
Materials and Methods

- <u>Cell Culture</u>: PBMCs were isolated from healthy donors by density gradient centrifugation (Ficol-Paque) supplied by the Blood Bank from healthy donors. Cells were cultured in RPMI media with 10% FBS, 1% L-glutamine, 1% penicillin/streptomycin, and 3 pg/ul recombinant human IL-2.
- *In-vitro* Treatments: Acute ethanol exposure was performed by incubation in media with 25mM or 50mM of ethanol for 24 hours. After ethanol exposure, cells were stimulated with 50 ng/mL phorbol myristate acetate (PMA) and 1 ug/mL lonomycin for 4 hours followed by RNA isolation.
- <u>RT-qPCR</u>: cDNA was then generated for each sample. Expression was measured using RT-qPCR on a Bio-Rad CFX96 Thermo Cycler using SSO Advanced Universal SYBR Green supermix for the following metabolism-related genes: Raptor, Rictor, and BCL-6. Expression was normalized to the housekeeping gene **RPS13**.
- Data Analysis: qPCR data was normalized to the untreated for each donor. All data was analyzed using Excel & GraphPad Prism 8.0. Statistical testing was performed using a 2-way ANOVA with Tukey's multiple comparison test. An alpha error < 0.05 was considered statistically significant.

PBMC expression of metabolic genes after acute 25 mM ethanol exposure.



PBMC expression of metabolic genes after acute 50 mM ethanol exposure.



were not statistically significant.

Conclusions

- We did not observe an ethanol effect.
- We saw a decrease in Rictor related to stimulation however there was no ethanol effect.
- There was no statistically significant difference in other genes based upon either stimulation or ethanol.

Future Experiments

- Continue PCR to determine gene expression for the alternate concentrations (25mM & 50mM) and timepoints (7 day).
- Assess differentiation of CD4 by flow cytometry after ethanol exposure

