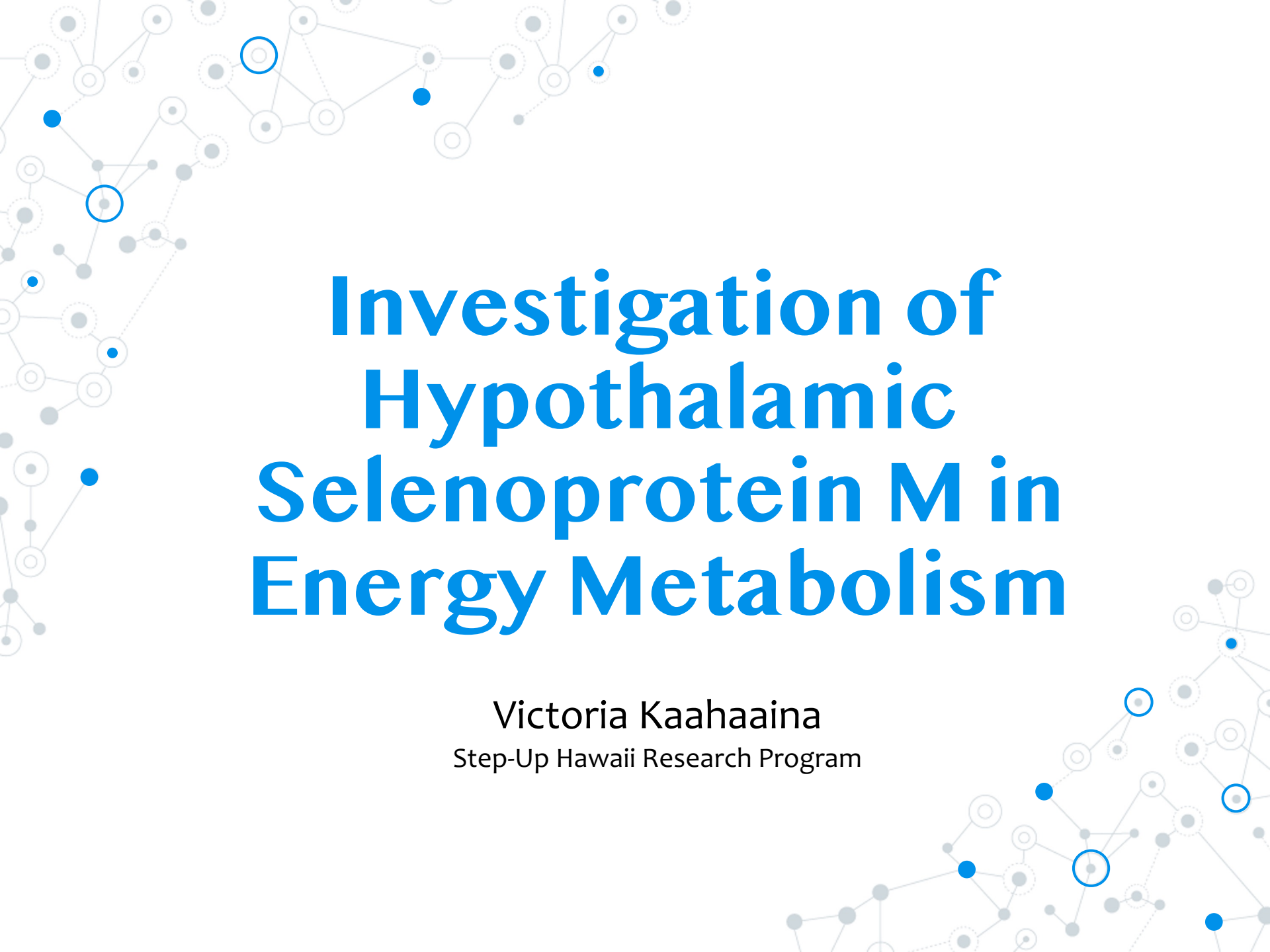




Investigation of Hypothalamic Selenoprotein M in Energy Metabolism

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Step-Up Hawaii Research Program

The background of the slide is a light gray network diagram. It consists of numerous small circles, some of which are solid gray and others are hollow with a gray outline. These circles are interconnected by a web of thin, light gray lines, creating a complex, interconnected pattern that resembles a molecular structure or a network graph.

Selenium and Selenoproteins

Selenium

- **Selenium (Se)** is an essential micronutrient
 - Important role in maintaining human health
 - Required for:
 - Normal thyroid hormone metabolism
 - Brain development
 - Immune function
 - Obtained primarily through diet
- Proteins with amino acid **Selenocysteine**
 - 25 genes in humans and 24 genes in mice encode selenoproteins

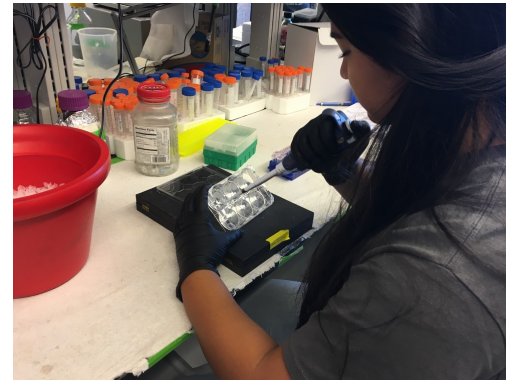
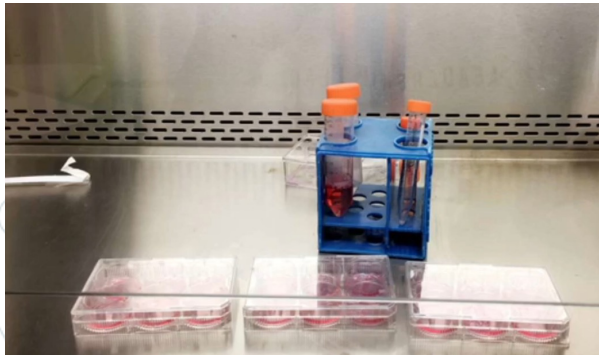
SelM

- A Thioredoxin-like protein
- Highly expressed in the brain
- Protects against oxidative stress



Last Year's Research

- **The Role of Selenium in Hypothalamic leptin signaling**
 - **Leptin**- Hormone released from fat that regulates energy homeostasis
 - **Leptin Resistance**- Failure of leptin to stimulate cell signaling pathways that promote energy metabolism
- **Hypothesis**- Selenium deficiency promotes hypothalamic leptin resistance
- **Experiment timeline**
 - Transfected cells, treated with Leptin, collected protein and analyzed by western blotting
- Results were inconclusive
 - SelM was significantly affected by the selenium treatment



The background of the slide is a light gray network diagram. It consists of numerous small circular nodes, some of which are solid gray and others are hollow with a gray outline. These nodes are interconnected by a web of thin, light gray lines, creating a complex, organic pattern that resembles a molecular structure or a data network. The overall aesthetic is clean and technical.

Experiment

SelM KO Mice

- Lack a functional SelM gene sequence
- SelM KO Mice developed leptin resistance and obesity at 18-20 weeks old when raised on a standard diet

Mice used in Hypothalamic Microarray Study

- 10- week-old males
- Similar body weight



Experimental Design

Microarray

Identify genes affected by SelM deletion



qPCR

Verify specific identified genes

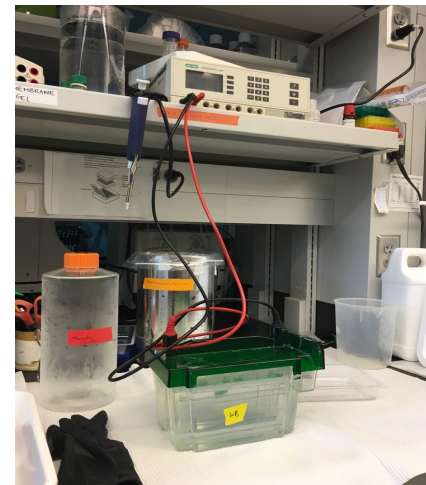
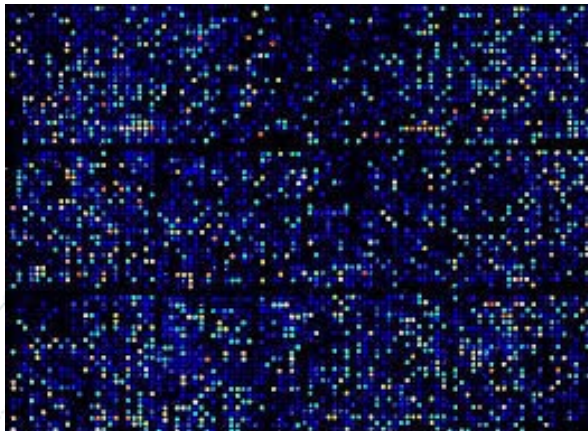
- p21
- PGC1 α
- TXNIP
- UCP2



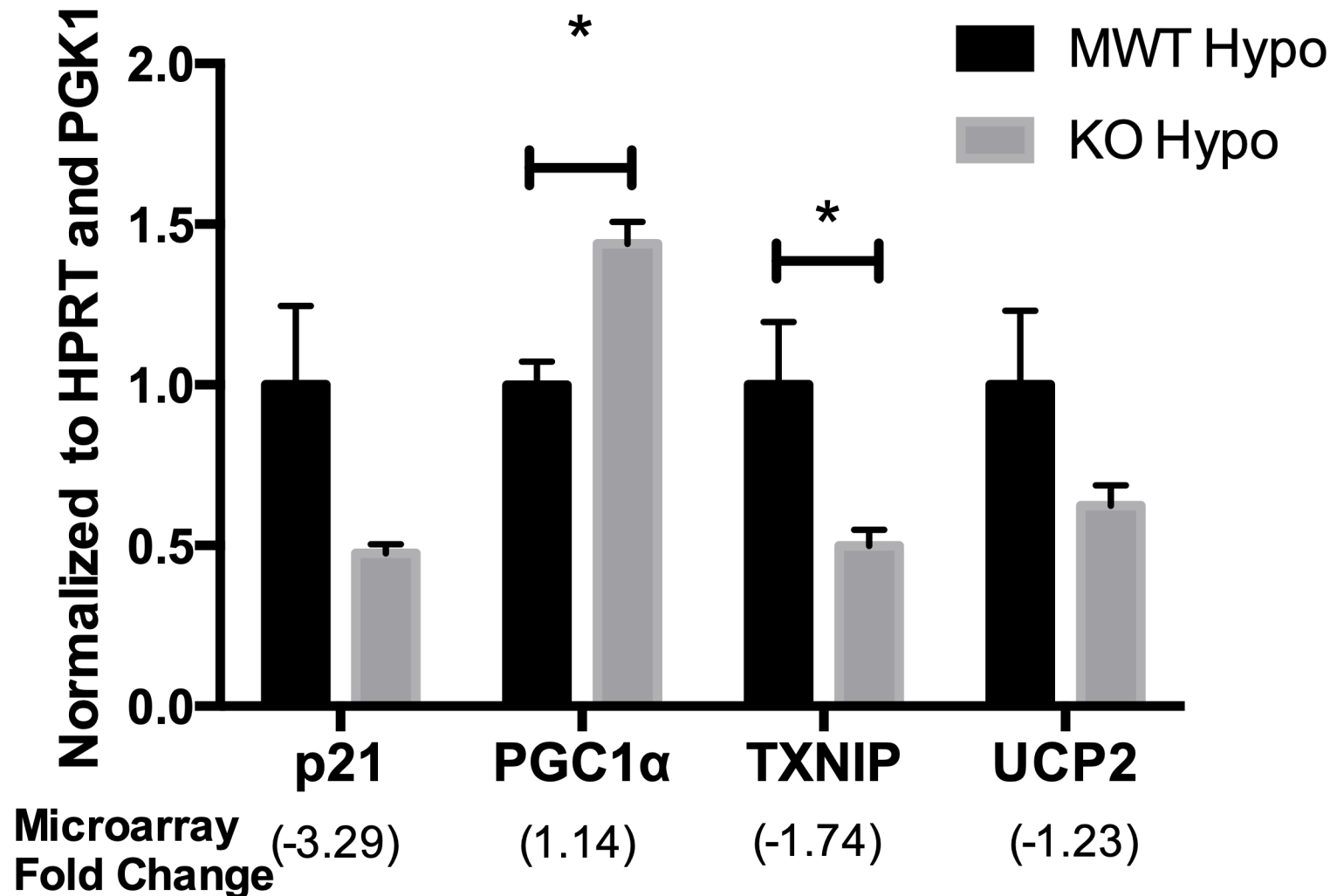
Western Blotting

Probe for genes verified in qPCR

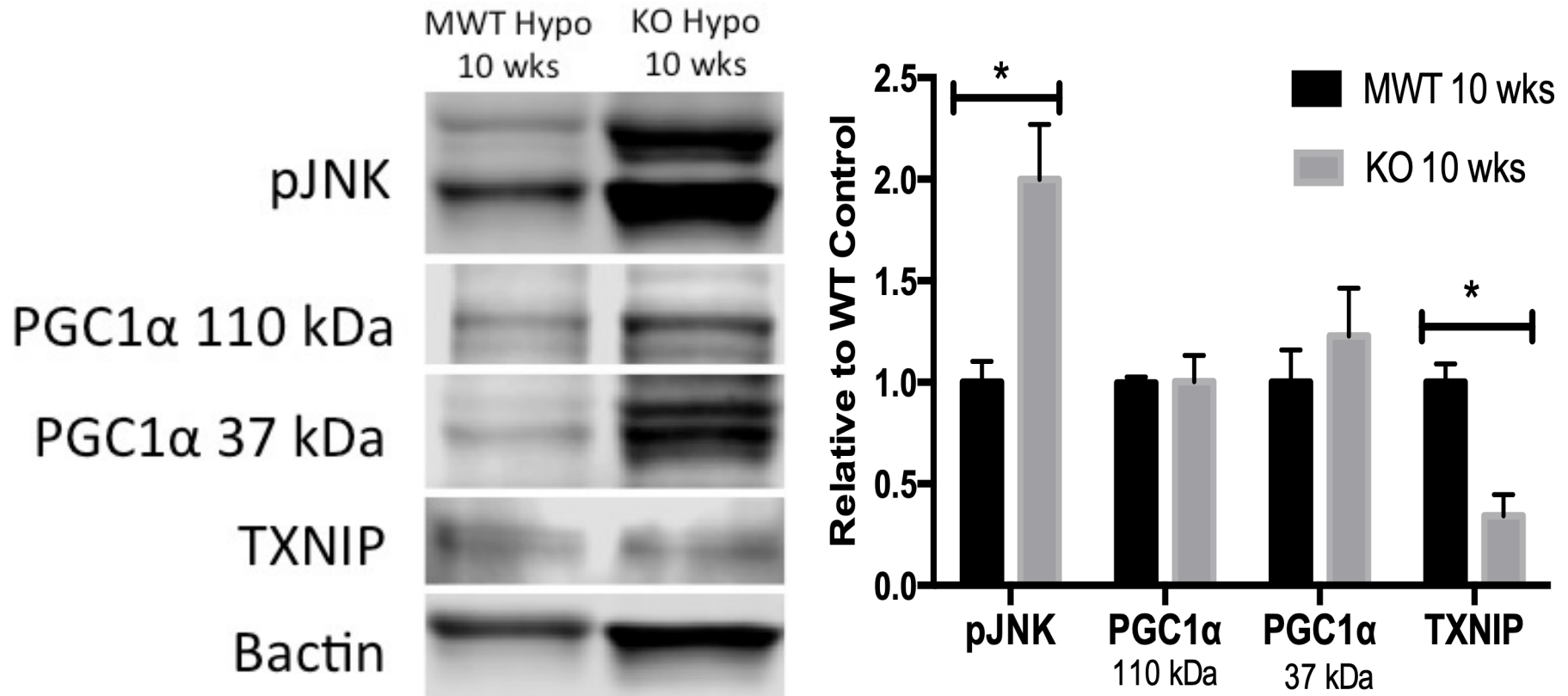
- PGC1 α
- TXNIP



qPCR Data: Hypothalamic Tissue

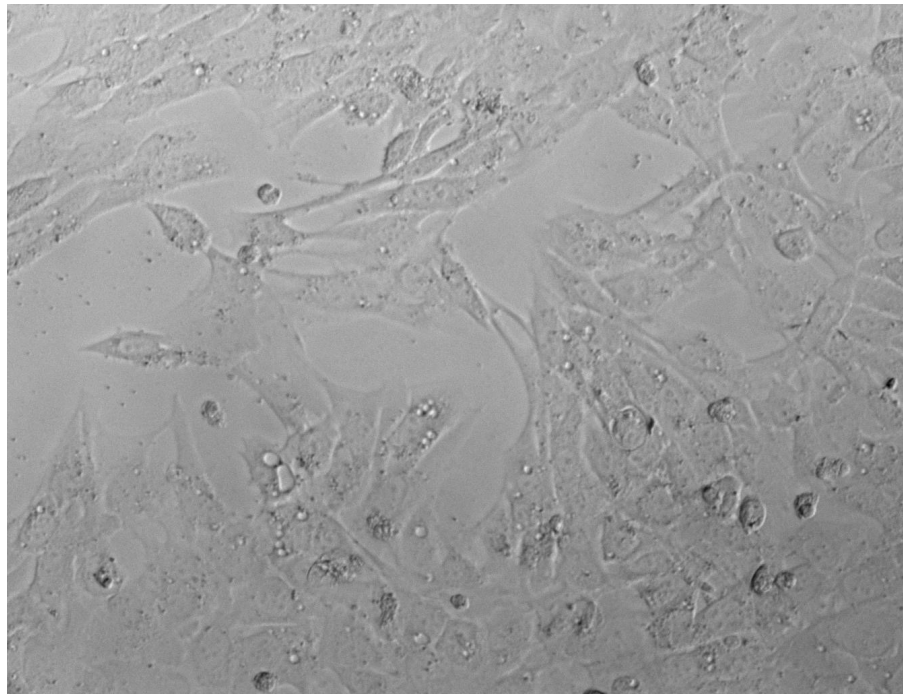


Western Blot Data: Hypothalamic Tissue

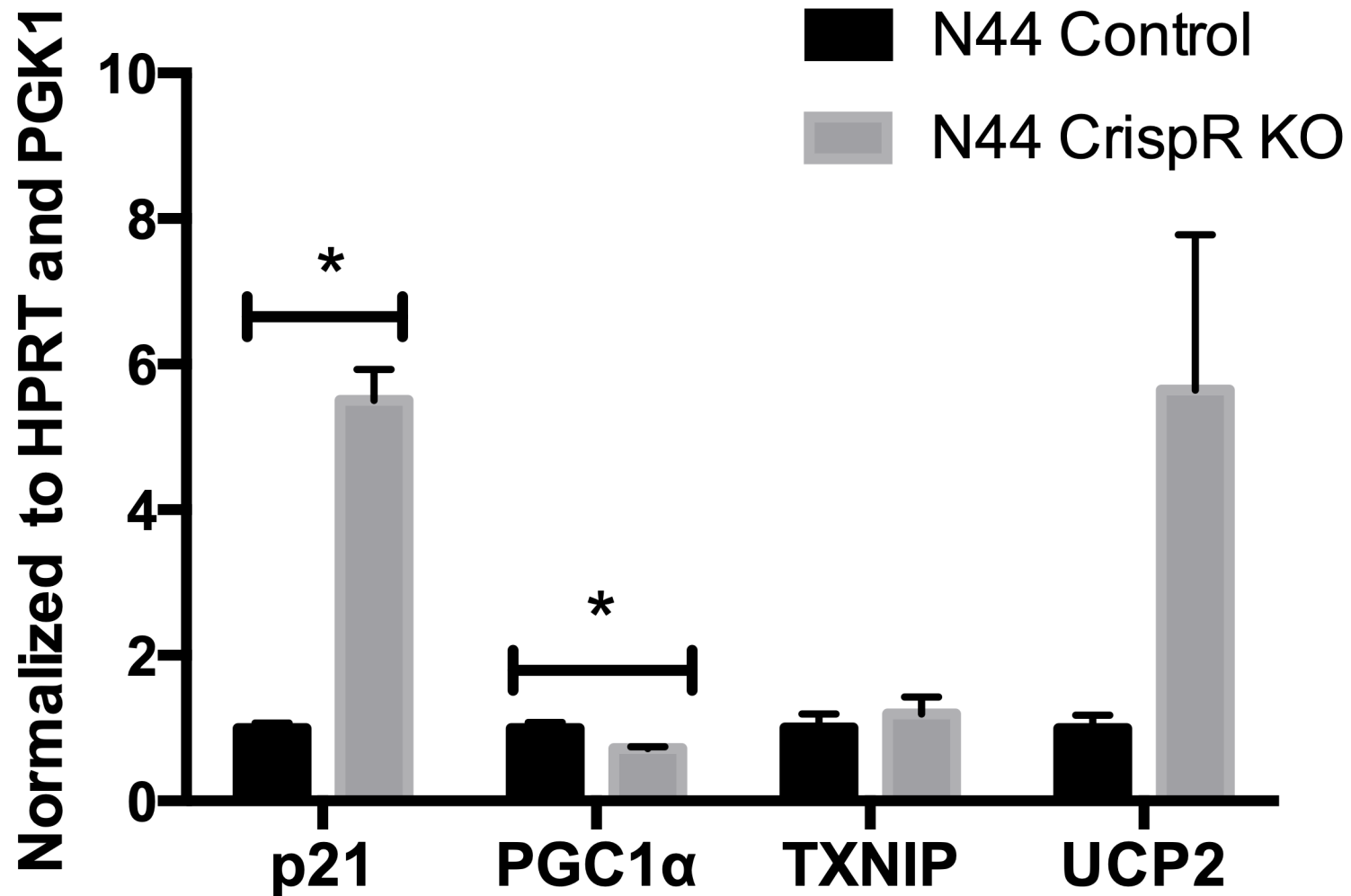


Cell line= mHypoE44

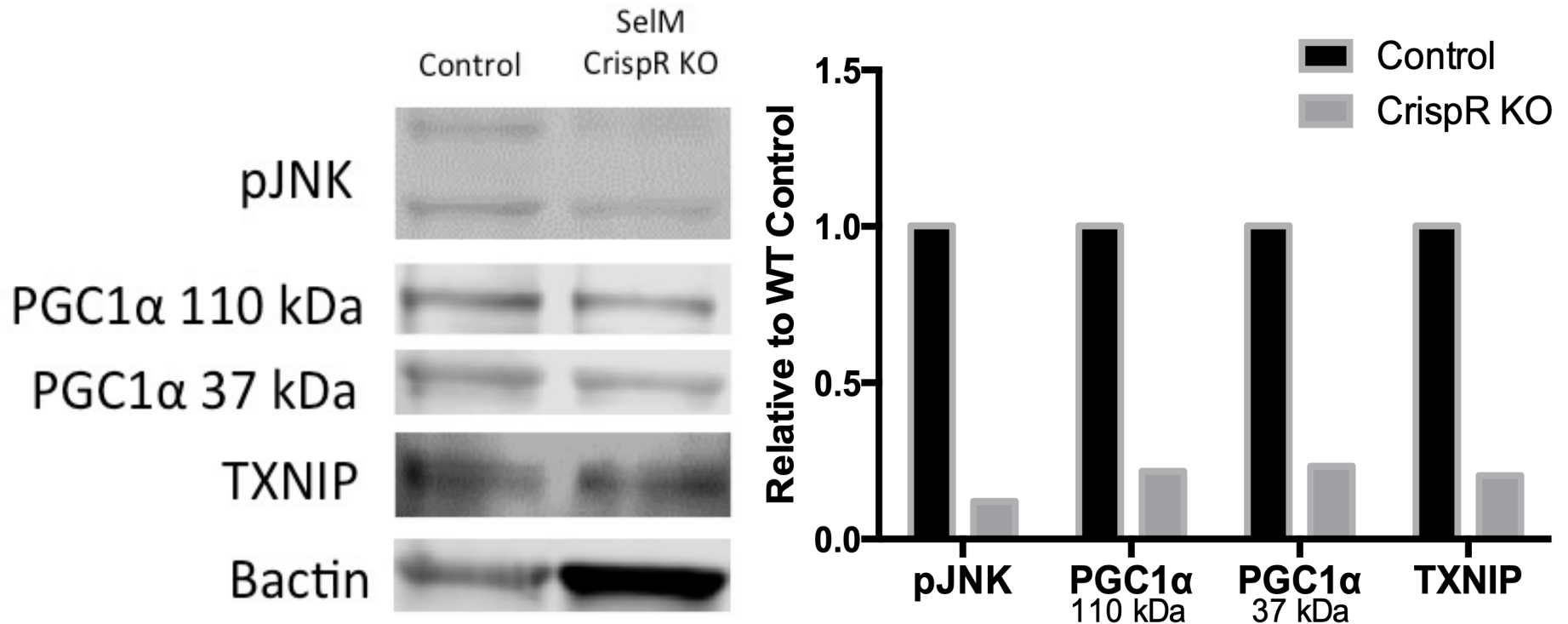
- Immortalized cell line derived from embryonic mouse hypothalamic neurons



qPCR N44 Immortalized Cell Line



Western Blot Data: N44 Immortalized Cell Line



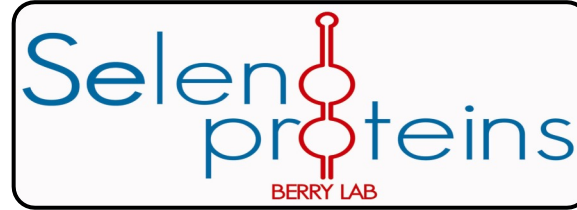
Conclusions

- SelM deletion significantly affects expression of p21, PGC1 α , and TXNIP
- Increased hypothalamic ER stress (pJNK levels) precedes disturbances in energy homeostasis previously reported in SelM KO mice



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Dedicated to Dioniso Domingo 1931-2017

