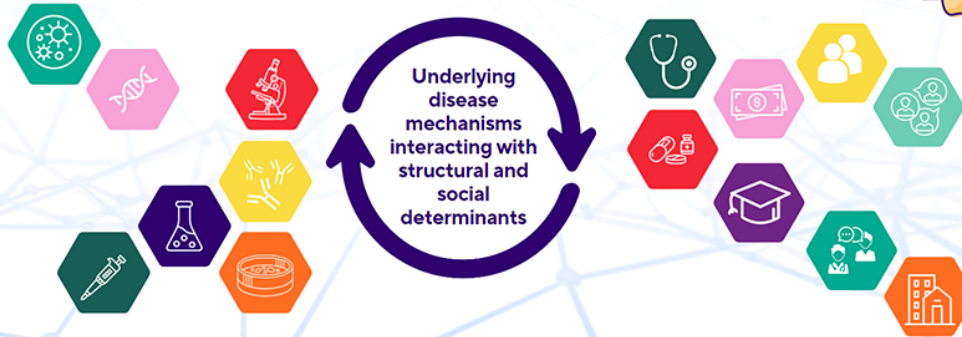




19TH ANNUAL CHILD HEALTH RESEARCH DAYS
Outcomes in Child Health



October 25 + 26, 2023 | RBC Convention Centre, Winnipeg, Manitoba

Abstract Submission Form

CHR D 2023: Abstract Submission Form

Submitter Name

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Presenter Name

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Presenter Status

PhD Student

Research Category

Basic Science

Role in the project

Design
Perform Experiments
Analyze Data
Write Abstract

Title

Gestational Diabetes Mellitus Induces Cardiac Dysfunction and Increases the Acetylation of Metabolic Enzymes in the Offspring Heart

Background

Exposure to gestational diabetes mellitus (GDM) increases risk of cardiovascular disease in offspring later in life. Previously, our lab found that GDM exposure impaired mitochondrial respiration by the cardiomyocytes of fetal offspring that was associated with cardiac dysfunction. Protein lysine acetylation is a post-translational modification that regulates activity of metabolic enzymes within the mitochondria.

Objective

To determine how GDM exposure affects protein lysine acetylation in the offspring heart.

Methods

GDM was induced by feeding female mice a high fat sucrose (HFS; 45% fat) diet for 6 weeks prior to mating and throughout pregnancy. Control lean dams were fed a low fat (LF; 10% fat) diet. Offspring from Lean and GDM dams were fed LF and HFS diets. Echocardiography was performed in 15-week-old offspring. Mitochondria were isolated from offspring hearts and acetylated peptides were extracted via immunoprecipitation and quantified by mass spectrometry.

Results

GDM and HFS diet induced cardiac hypertrophy (Lean-LF vs. GDM-HFS $p=0.015$) and diastolic dysfunction (Lean-HFS vs GDM-HFS $p=0.006$) in 15-week aged offspring. GDM exposure differentially

altered the acetylation of mitochondrial peptides, which was exacerbated by a postnatal HFS diet (GDM-HFS vs Lean-LF 88 peptides, $p < 0.05$). Functional classification revealed prominent representation of acetylated enzymes involved in the TCA cycle, fatty acid oxidation, respiratory electron transport, and mitochondrial biogenesis pathways in the hearts of GDM offspring.

Conclusion

GDM-induced cardiac hypertrophy and diastolic dysfunction in the offspring. These impairments in cardiac structure and functions were found in tandem with increased acetylation of cardiac mitochondrial enzymes regulation energy metabolism. Cardiac mitochondrial enzyme acetylation represents a novel molecular mechanism that contributes to GDM-induced mitochondrial dysfunction and cardiomyopathy in the offspring.

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