

# Epigenetics for Breast Cancer Prevention: A Family Case-Control Study

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# **BACKGROUND**

- Breast cancer most common malignancy and second leading cause of cancer death among women.
- Methylenetetrahydrofolate reductase (MTHFR) one of the most studied gene variations associated with breast cancer affecting epigenetic pathways.
- Epigenetic (gene-environment interaction) modifications from unhealthy environment and behaviors can lead to cancer.

# **PROBLEM STATEMENT**

- Study results are inconsistent on breast cancer gene variations as risk factors for breast cancer.
- Inconsistencies are due to environmental and behavioral factors, i. e. folate, alcohol intake, & smoking, that warrant further investigations.



# **PURPOSE**

• Investigate the associations of epigenetic risks for breast cancer.

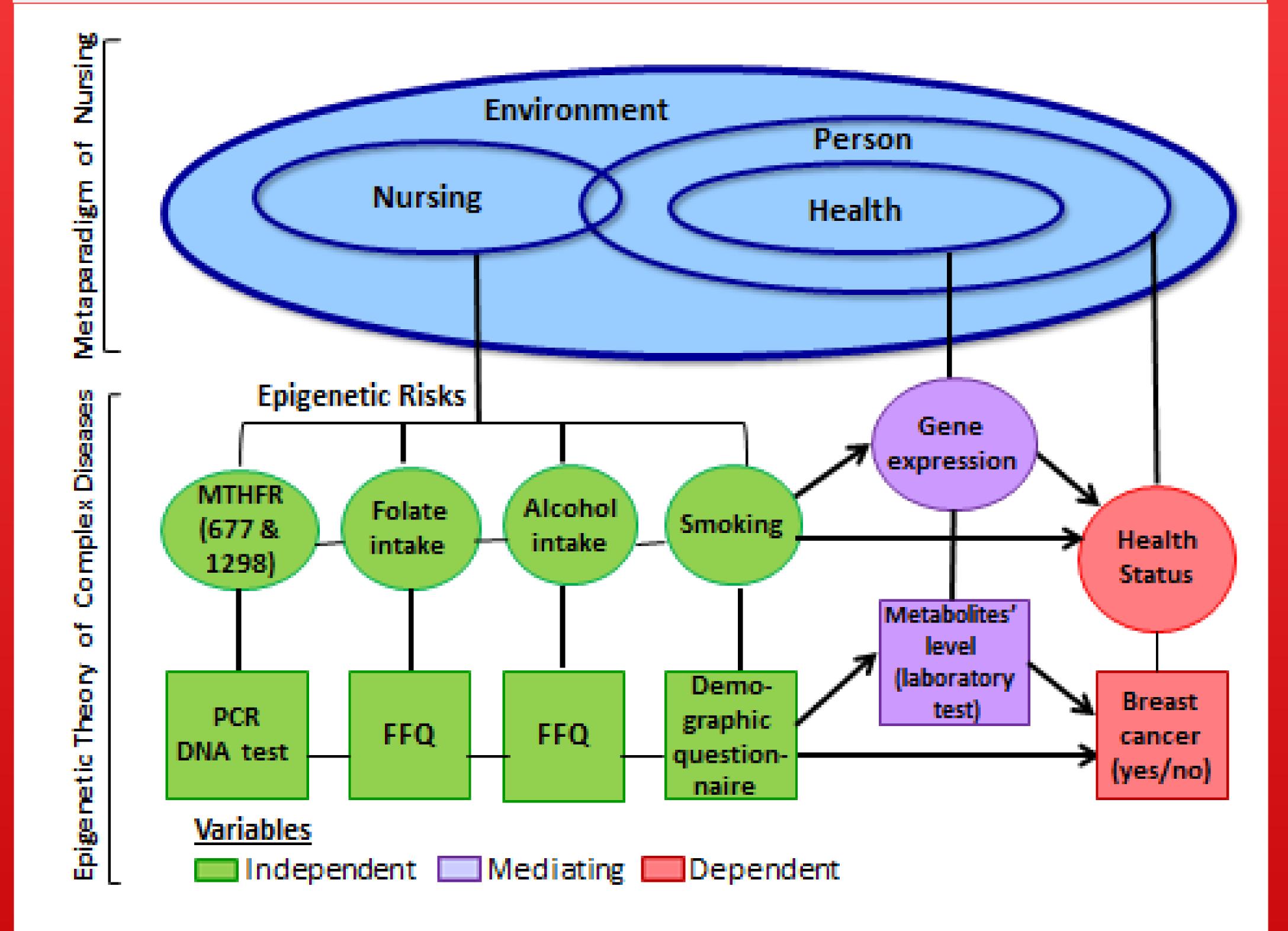
# **SIGNIFICANCE**

 Identifying health behaviors offers opportunity for future interventions that target epigenetic pathways for breast cancer prevention.

# CONCEPTUAL THEORETICAL FRAMEWORK:

Nursing Metaparadigm and Epigenetic Theory of Complex Diseases

Cancer is a complex disease linked to epigenetic misregulation affected by endogenous and exogenous factors.



MTHFR Cytogenetic
Location: 1p36.3
Chromosome 1

# **METHODS**

# **SETTING**

• Southern California counties – Los Angeles, San Bernardino, Riverside, and Orange

# **SAMPLE**

 Total of 60 female participants - 30 breast cancer cases and one of their family members as controls will be recruited from California Cancer Registry.

### **INSTRUMENTS**

 Food Frequency Questionnaire (FFQ), 2000 Cancer Module National Health Interview Survey, and Demographic Questionnaire

# **ANALYSIS**

# LABORATORY ANALYSIS

 Salivary and/or blood samples for MTHFR gene variations (SNPs 677 1298) using PCR DNA test, serum folate, and metabolites' analyses

# STATISTICAL ANALYSIS

- Pooled analysis of relative risks to determine rate differences between cases and control groups
- Correlation tests, t-test, and chi-square to determine associations among variables

**DATA COLLECTION** - in progress

# CONCLUSION

 Associations of epigenetic risks with breast cancer will serve as foundation for future development of innovative behavioral interventions; personalized healthcare for breast cancer prevention and control based on genome health.

# THEORETICAL IMPLICATIONS

 Foundation for new hypotheses and possible generation of middle range theory for epigenetics of cancer.

# **CLINICAL IMPLICATIONS**

- Significant collaborative role of Nursing in translating knowledge of epigenetics into health promotion and disease prevention.
- Evolution of new targeted approach in diagnostics and treatment management for breast cancer.

# HEALTH POLICY IMPLICATIONS

• Impact for high quality healthcare with enhanced genome health awareness and competencies among healthcare professionals.

# **FUTURE RESEARCH**

- Future studies to establish epigenetic risks that can be reversed to prevent cancer.
- Prospective development of personalized behavioral intervention for cancer prevention and control.

# **ACKNOWLEDGEMENT**

Strength and perseverance are nurtured within. It is an honor & privilege to be supported & guided by inspiring members of my Dissertation Committee.



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