# Consequences of folate depletion during development for DNA methylation and gene expression in the fetal mouse

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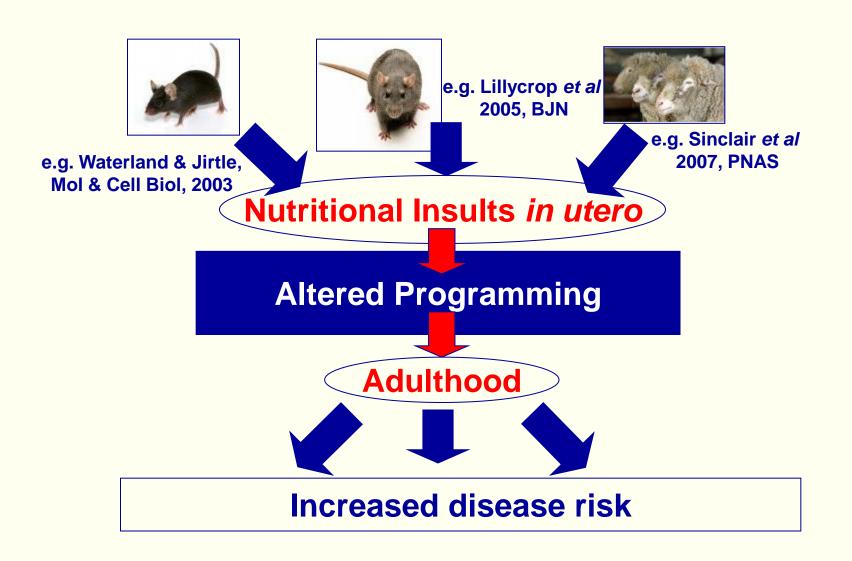
<sup>2</sup>BiGCaT, Maastricht University

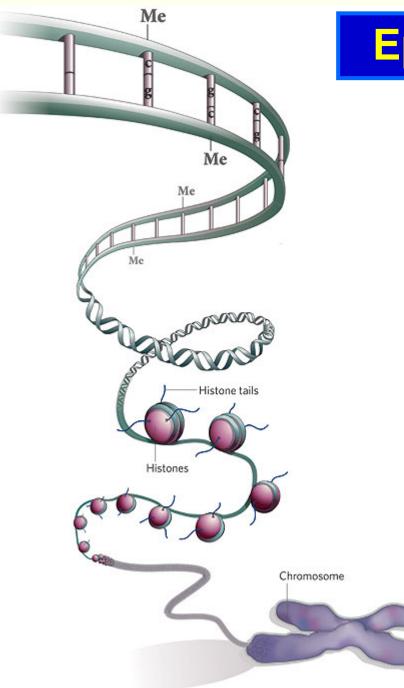






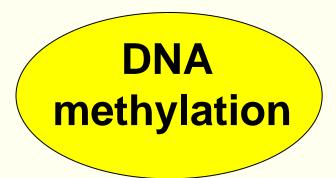
# Developmental Origins of Health & Disease





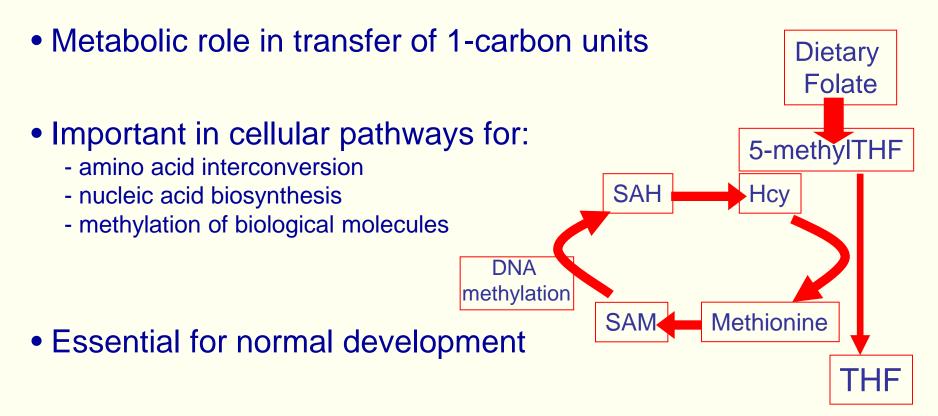
# **Epigenetic Mechanisms**

- Candidate mechanisms for developmental programming
- Established in utero
- Modifiable by environmental factors
- Influence gene expression

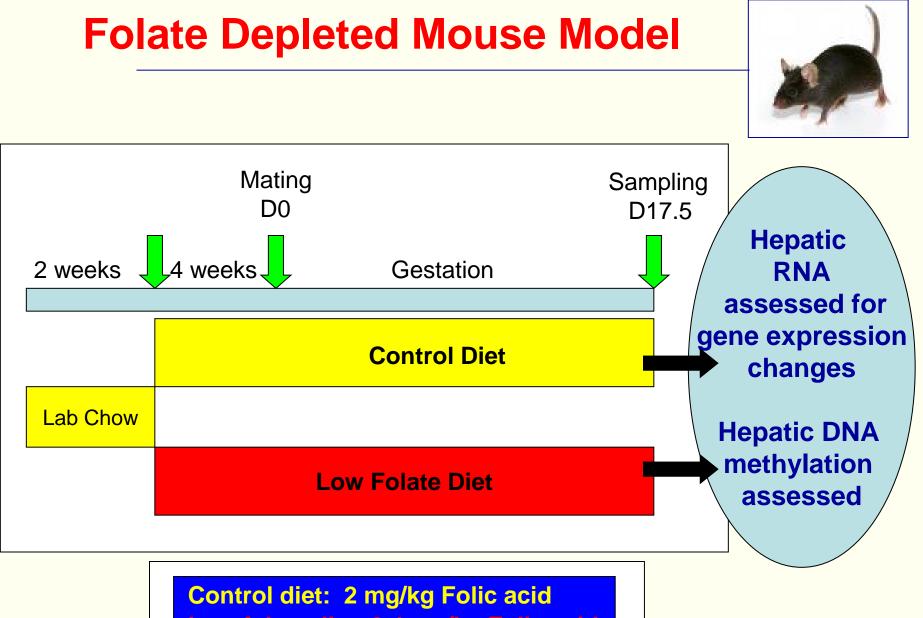


Qiu J (2006) Nature 441, 143-145

#### Folate

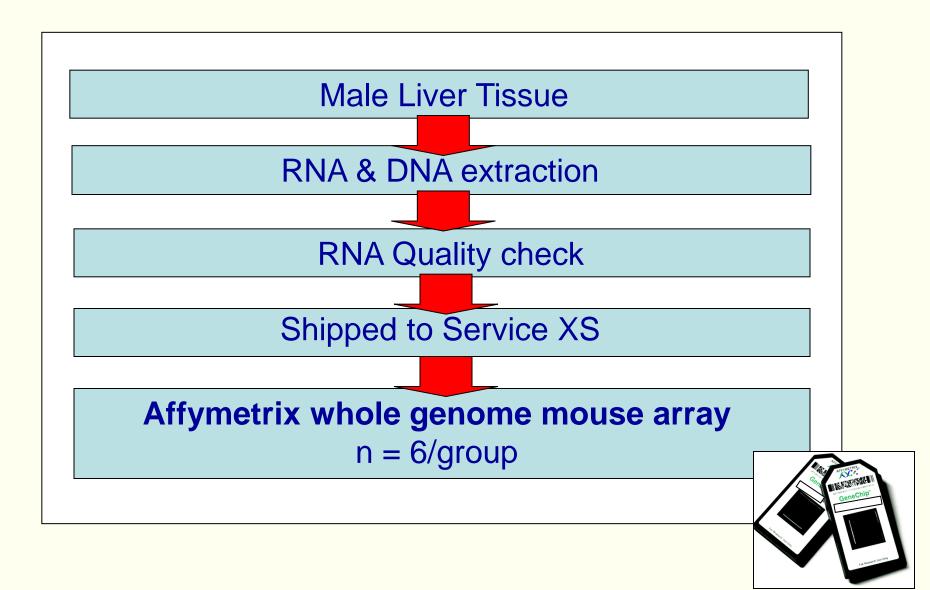


 Low folate intake during pregnancy is associated with increased risk of NTDs Hypothesis Offspring of folate depleted mothers have altered gene expression caused via DNA methylation changes

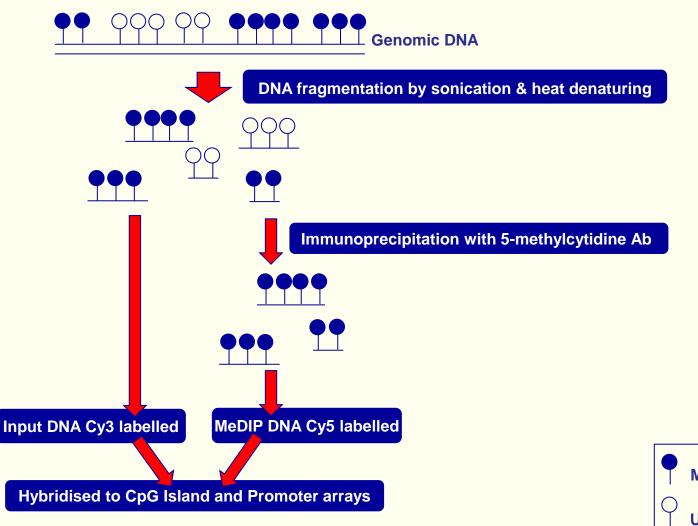


Low folate diet: 0.4 mg/kg Folic acid

#### **Gene Expression Arrays**



# Methylated DNA Immunoprecipitation (MeDIP) for Methylation Arrays



Methylated CpG

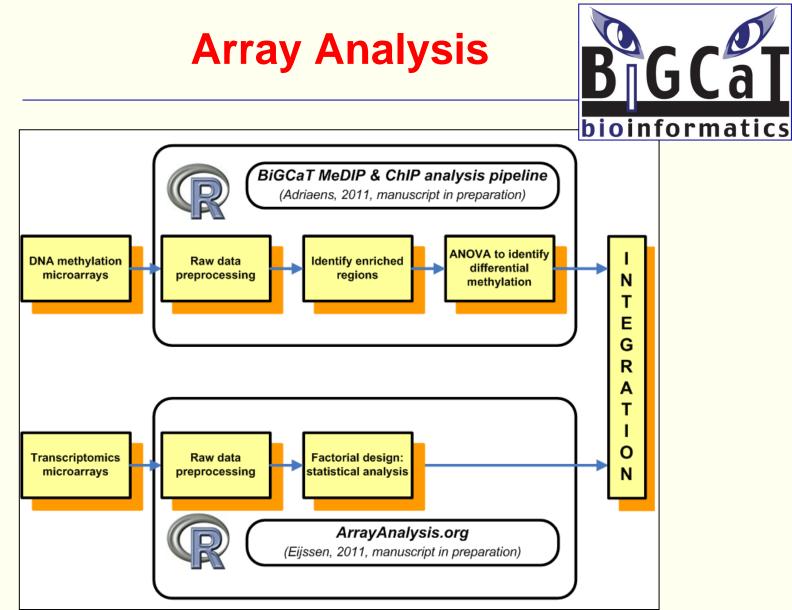


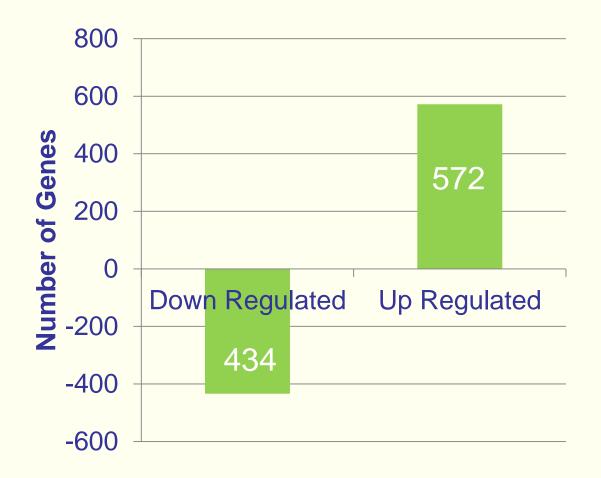
Figure composed by <u>Michiel Adriaens</u>

Poster: 'Consequences of folate depletion during development and

high fat intake from weaning on adiposity, gene expression and DNA methylation in adult mice. "

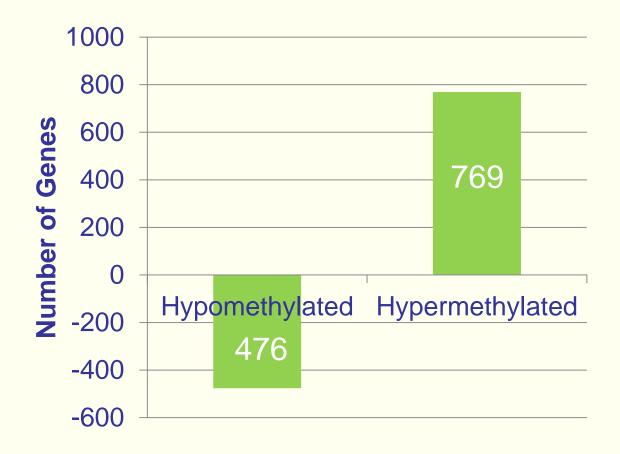
# Gene expression changes in response to low maternal folate

- Fold change
  >1.2 or < -1.2</li>
- Significant
  P value <0.05</li>

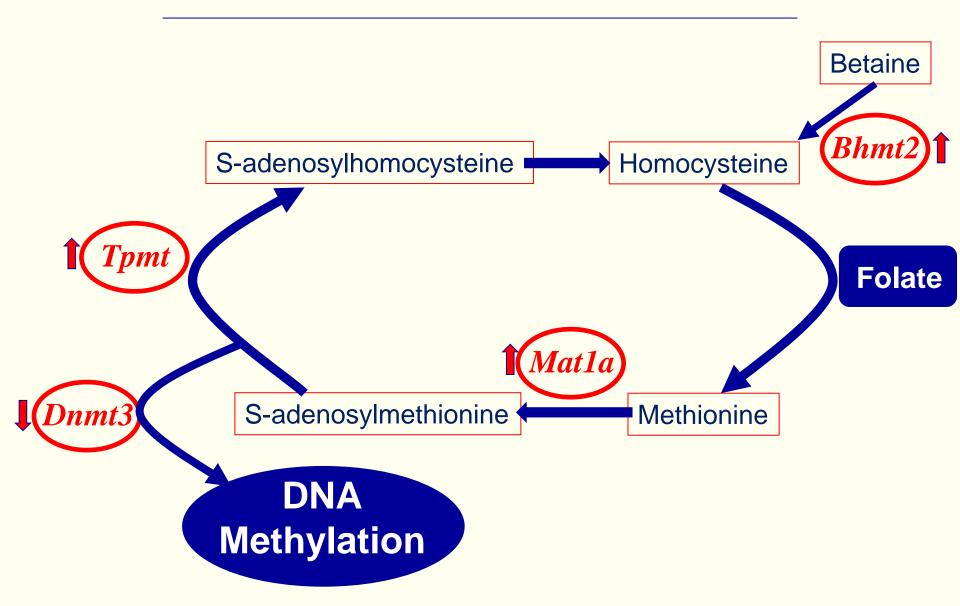


# DNA methylation changes in response to low maternal folate

Significant
 P value <0.05</li>

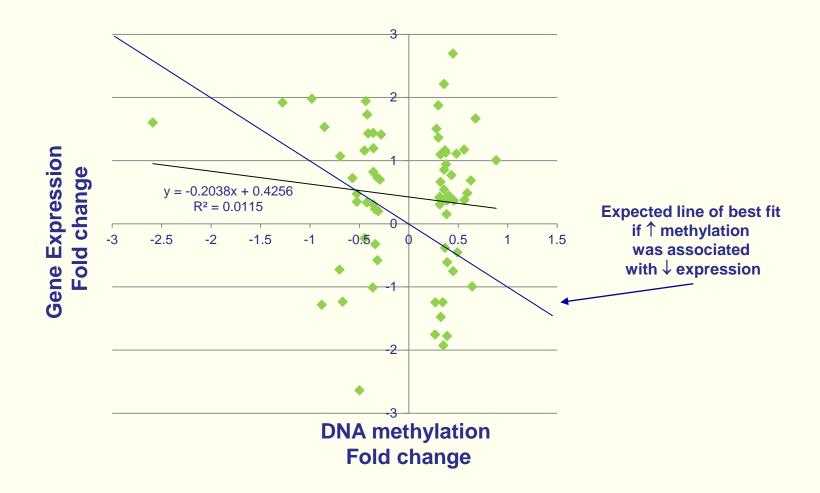


# **Possible Mechanism**



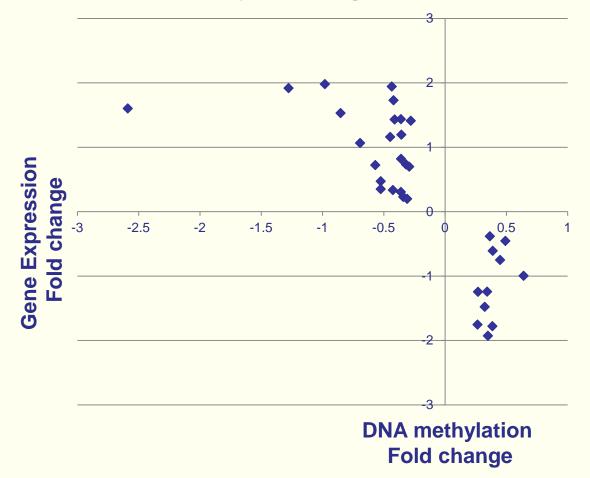
# Genes with altered expression & DNA methylation

72 genes were differentially expressed AND differentially methylated in response to low maternal folate



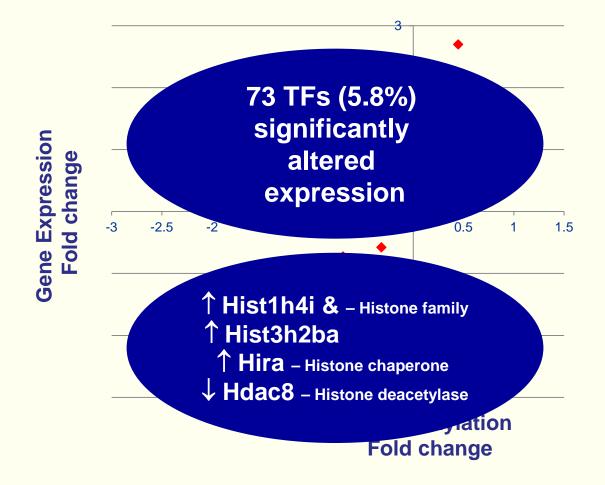
#### **Direct Effects**

Changes in gene expression of 33 genes were potentially caused by changes in DNA methylation: DNA methylation ↑, gene transcription↓



#### **Indirect Effects**

Changes in gene expression of 39 genes were caused by other mechanisms: altered expression of transcription factors, histone modifications or post-transcription µRNA events



### **Conclusions**

- Maternal low folate alters both gene expression & DNA methylation in fetal liver
- More genes were hypermethylated than hypomethylated in response to low folate
- 72 genes were differentially expressed AND methylated in response low maternal folate
- Suggests mechanisms other than DNA methylation are involved in observed altered gene expression e.g. changes in transcription factor expression, histone modifications etc
- It is essential to consider both direct and indirect effects when integrating epigenomic and transcriptomic data

# **Acknowledgments**

# **Technical Support**





Adele Kitching Sofia Lisanti

# **Collaborators**



Chris Evelo Michiel Adriaens

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