# Transgenerational Nutrition

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# Overview of talk

- Research Program
- Observations Ca/P conditioning
- Long term impacts of Environment (Diet)
- Epigenetics
- One Carbon Metabolism Feeding the machinery
- Challenges in Poultry Production
- Can we feed the next generation(s)?





### In life there is no such thing as a free lunch.



# You can only cut the pie (genetic resources) into so many pieces.



Ultimately the "pie" is finite, there is a limit to the genetic potential of an organism.



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#### The practical problem: Manure Phosphorus









### Long term impact of dietary conditioning





### Epidemiological Observations: Fetal origin of Adult Disease- The Barker Hypothesis

 1989 David Barker found in inverse relationship between birth weight and death from heart disease in UK.



- Studies confirmed by "Dutch Hunger Winter" when food was cut off in occupied Netherlands. Individuals born during this time, and their offspring have high incidence of insulinresistance as adults.
- "Fetal Origin of Adult Disease" confirmed for:
  - Coronary heart disease
  - Hypertension
  - Type II diabetes
  - Metabolic disorders, ect...

## **Epigenetics and Diet-Honeybee**



- Larva destined to become a queen is fed large quantities of royal jelly by worker bees (nurse)
- Royal Jelly consumed by queen larva inactivates the Dnmt3 gene so genes can remain active and result in queen characteristics
- When Dnmt3 is active it silences genes in bee larva and inhibits them from developing into a queen
- Effects morphology, physiology, life span and behavior
  - Egg laying
  - Kills rival queens
  - Produce communication sounds
  - Pheromone production
  - Mating flights







## Transgenerational Epigenetics



Gregarious

Solitarious

Schistocerca gregaria locust (left) and Schistocerca gregaria grasshopper (right).

The hind legs of the locust are set at a different pitch. The wings are smaller while the brain is larger, and yet these two insects are the same species.

The grasshopper can transform into a locust in a matter of hours with the right stimulus (hormones, physical or visual stimulation), and will maintain this phenotype for 3-4 generations if returned to low density.



### Dietary impact on DNA methylation status



**Figure 1.** Methylation status of CpG sites within the A<sup>vy</sup> IAP in genetically identical A<sup>vy</sup>/a littermates. (A) A contraoriented IAP insertion within pseudoexon 1A (PS1A) of the murine Agouti gene. A cryptic promoter (short arrow labeled A<sup>vy</sup>) drives ectopic Agouti expression. CpG sites 1–9 are oriented in the 3' to 5' direction with respect to the IAP insertion, as shown. Transcription of A and a alleles initiates from a hair-cycle-specific promoter in exon 2 (short arrow labeled A,a). (B) Pseudoagouti animals exhibit the highest degree of CpG methylation at sites 4–9. Bisulfite sequencing reveals increasing intensity of the cytosine lane at CpG sites 4–9 within the A<sup>vy</sup> IAP in genetically identical A<sup>vy</sup>/a animals representing the five coat classes. (C) Genetically identical week-15 A<sup>vy</sup>/a mouse littermates representing the five coat-color phenotypes.

## Transgenerational Nutrition



### Environmentally induced epigenetic transgenerational inheritance

#### **Environmental toxicants**

Agricultural fungicides (Vinclozolin) Agricultural pesticides (Methoxychlor) Industrial contaminants (Dioxin/TCDD) BPA and phthalates (Plastic compounds) Herbicides (Atrazine and glyphosate)

#### Other types of exposure

Nutrition (High fat or caloric restriction) Temperature and drought (Plant health and flowering)





Flies

Plants

Worms



Fish

Industrial toxicants and biocides (Tributyltin) Hydrocarbons (Jet fuel JP8) Heavy metals (Mercury)

Insect repellants (Permethrin and DEET)

Smoking and alcohol Stress and trauma (behavioral)



Birds

Pesticides (DDT)







Humans

King and Skinner, 2020

Rodents



## Looking for Transgenerational evidence

- Evaluate dietary methyl donor effects
  - Reproductive and Growth Performance
  - Gene Expression
  - Individual gene methylation patterns
  - Epigenetic effects of diet on progeny performance across multiple generations

PhD – Chelsea Phillips



## Dietary Methyl donors

- <u>Choline</u>: Oxidized to betaine as a source of methyl groups in forming methionine from homocystine *McGinnis et al. (1944)*
- **<u>Betaine</u>**: Methyl Donor
- **<u>B12</u>**: coenzyme of methionine synthase which catalyz
- **Folic Acid**: carrier of single carbon units as 5-methyl THF which is donated to homocystine for the formation of methionine *McKay et al. (2011)*
- **<u>B6</u>**: Coenzyme to hydroxymethyltransferase which converts THF to 5,10-methylene THF
- Zinc: cofactor required for amino acid metabolism and DNA modification



## Nutrient Specifications

Methylation package (10X)	%	mg/kg
Choline Chloride 98%	0.5	7030
Betaine , 99%	0.5	5
B12, 98%	0.0015	1.5
Folic acid 95%	0.075	7.5
B6 (Pyridoxine 98%)	0.012	12
Zn Sulfate 99%	0.00987	99
Total methylation	1.09837	

Starter Nutrients	Amount
Crude Protein (%)	27.9
Met. Energy (KCAL/KG)	2,850
Calcium (%)	1.00
Avail. Phos. (%)	0.50
Met, total (%)	0.7392

Layer Nutrient	Amount
Crude Protein (%)	22.6
Met. Energy (KCAL/KG)	2,800
Calcium (%)	2.6
Avail. Phos. (%)	0.45
Met, total (%)	0.64

## **Trial Parameters**

#### **Phenotypes Collected for Each Generation**

<b>G0</b> Parents	BW	# of Eggs	Egg Wt	Body Composition	DNA Methylation Seq Data, RNAseq
<b>G1</b> Progeny	BW	# of Eggs	Egg Wt	Body Composition	DNA Methylation Seq Data, RNAseq
<b>G2</b> Progeny	BW	# of Eggs	Egg Wt	Body Composition	DNA Methylation Seq Data, RNAseq

Phillips and Ashwell, unpublished

## **Trial Parameters**

Significantly Different Phenotypes Collected for Each Generation				
<b>GO</b> Parents	<b>21d BW</b> p<0.0001	<b>Egg Wt</b> p<0.0001	<b>Liver Wt as % BW</b> p<0.0028	
<b>G1</b> Progeny	<b>Hatch BW</b> p<0.0001	<b>Egg Wt</b> p<0.0001		
<b>G2</b> Progeny		<b>Egg Wt</b> p<0.0001		

Phillips and Ashwell, unpublished





## Classification of CpG changes resulting from diet – G0



- 1,232,624 CpGs had sufficient reads to make a statistical inference
- Average reads per CpG site 39.3
- ~1.8 million CpGs predicted in chicken, ~70% coverage
- 98.6% of CpG sites are unchanged
- Pathway analysis of the ΔMeth loci Ingenuity

Phillips and Ashwell, unpublished



### Antigen Presenting Pathway Hypomethylation & Hypermethylation

#### MHC Class II Molecules

- Antigen Presenting Cells
- Initiate Immune Response

Common on antigen presenting cells but can be induced on other cells by interferon  $\gamma$ 

MHC II expression is regulated by CIITA

INF  $\gamma$  triggers the expression of CIITA

Triggered immune response

- Inflammation, swelling
- Possible B Cell activation

## Insights....and future

- Diet impacts offspring for at least 2 generations
- Current industry practice must effect commercial broilers
- Investigate loci associated with egg size and determine if the methylation pattern correlates with the phenotypes
- Challenge or stress studies?
- Treatment effect in a single sex?
- Current industry practice must effect commercial broilers

#### Pedigree

#### 1 female

These are the elite birds that produce future generations. At this stage, genetic selections are performed by the research & revelopment team at our pedigree farms.

#### Grandparents

#### 725 females

GPs are pure line birds that are placed into mixed-line mating. These birds live at Cobb farms or distributor farms and produce our "parent" products.



#### Great Grandparent

23 females

These pure line birds live at company-owned facilities. Phenotypic selections are done for males only. The eggs from the females are hatched to produce grandparent birds.



29,000 females



## Evaluate the impact of Precision Feeding



Breeders fed to increased body weights increases FCR in broiler progeny

Investigating the impact of graded feeding on breeder and broiler progeny gene expression patterns – altered stress pathways

Impact on DNA methylation?

Collaboration with Martin Zuidhof- University of Alberta



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### **Questions?**