

# Early-programming of dairy cattle, a potential explanation to the adaptation to climate change

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# Context: Epigenetics

**FEEDING**



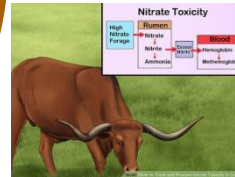
**HEAT STRESS**



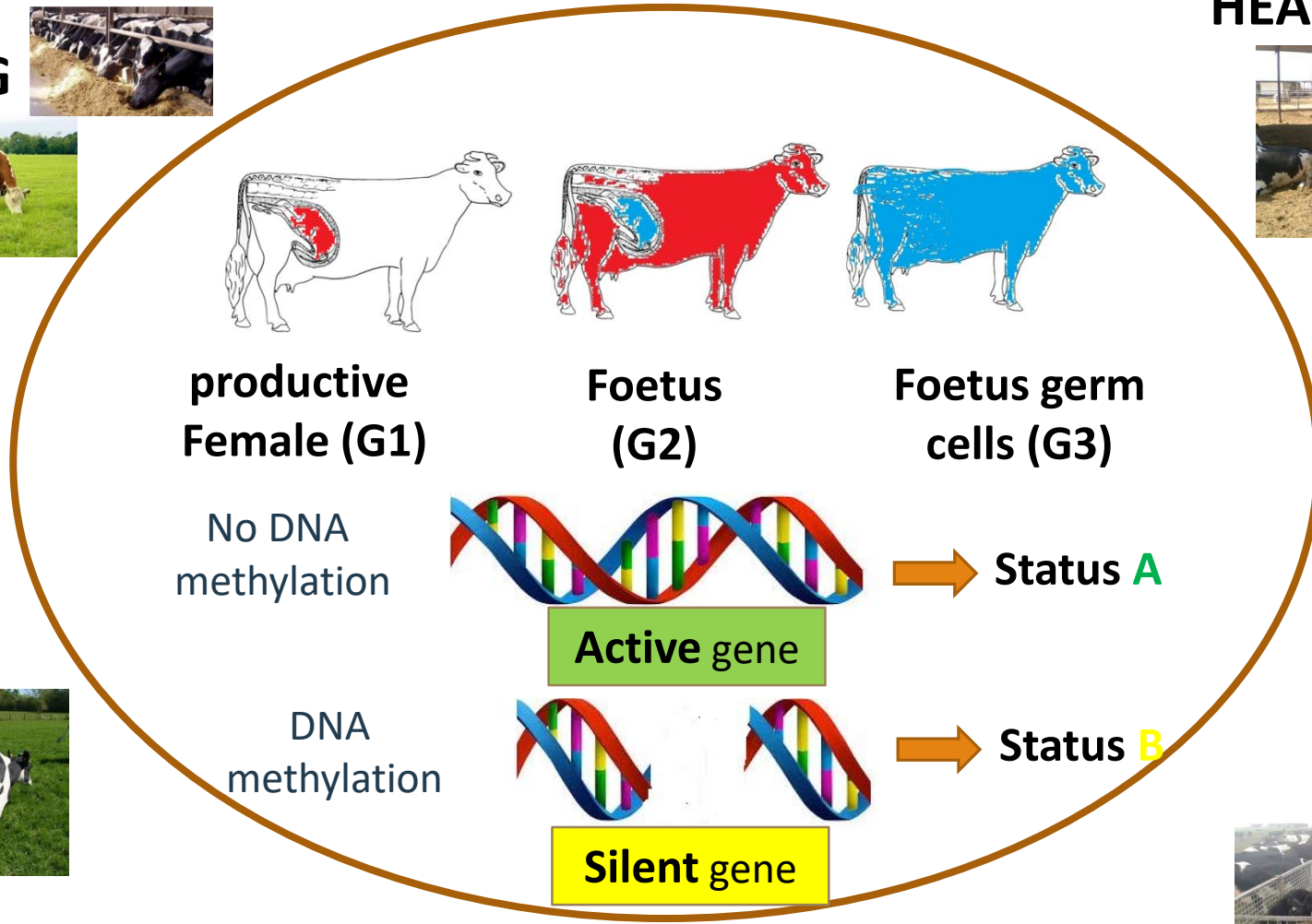
**HEALTH**



**TOXICITY**

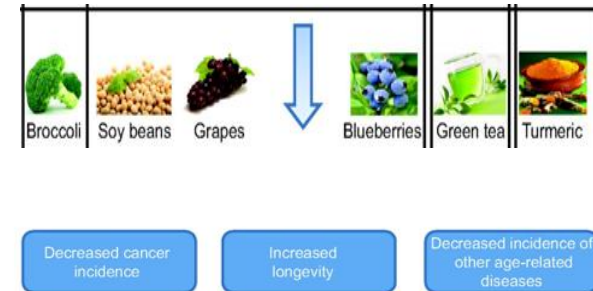


**BARN**



# Opportunities in dairy cattle

- Human disease and prevention (e.g. cancer)
  - Genetic make-up of disease development (or not)
  - Early programming (e.g. epigenetic diets)



Michael Daniel, and Trygve O. Tollefsbol J Exp Biol 2015;218:59-70

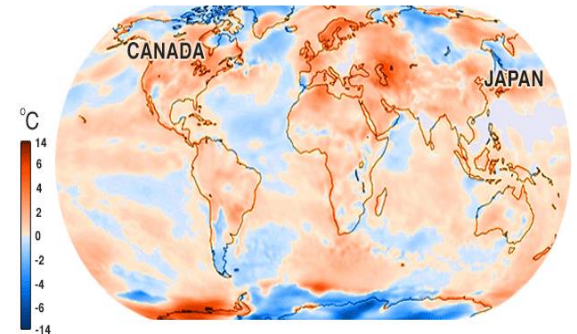
- Those studies can be explored dairy cattle:
  - Identify practices associated to epigenetic changes
  - Detect genotypes with reduced susceptibility to epigenetics



**G x E & Epi-Genome Wide Studies**

# Challenges: adaptation to climate

Where in the world temperatures are above and below average for 1-20 July



Temperatures are measured against the average for the period 1981-2010  
Source: Copernicus Climate Change Service, European Centre for Medium-Range Weather Forecasts **BBC**

- Heat stress (HS):
  - Temperate regions

- Recent studies on expose of dairy cows to HS during the conception period **PLOS ONE**



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**Season of conception is associated with future survival, fertility, and milk yield of Holstein cows**

P. J. Pinedo\*<sup>1</sup> and A. De Vries†



RESEARCH ARTICLE

Periconceptual Heat Stress of Holstein Dams Is Associated with Differences in Daughter Milk Production and Composition during Multiple Lactations

Britni M. Brown<sup>1</sup>, Jon W. Stallings<sup>2</sup>, John S. Clay<sup>3</sup>, Michelle L. Rhoads<sup>4\*</sup>

**Evidence that season of conception affects offspring lifetime**

- Mainly at the phenotypic level

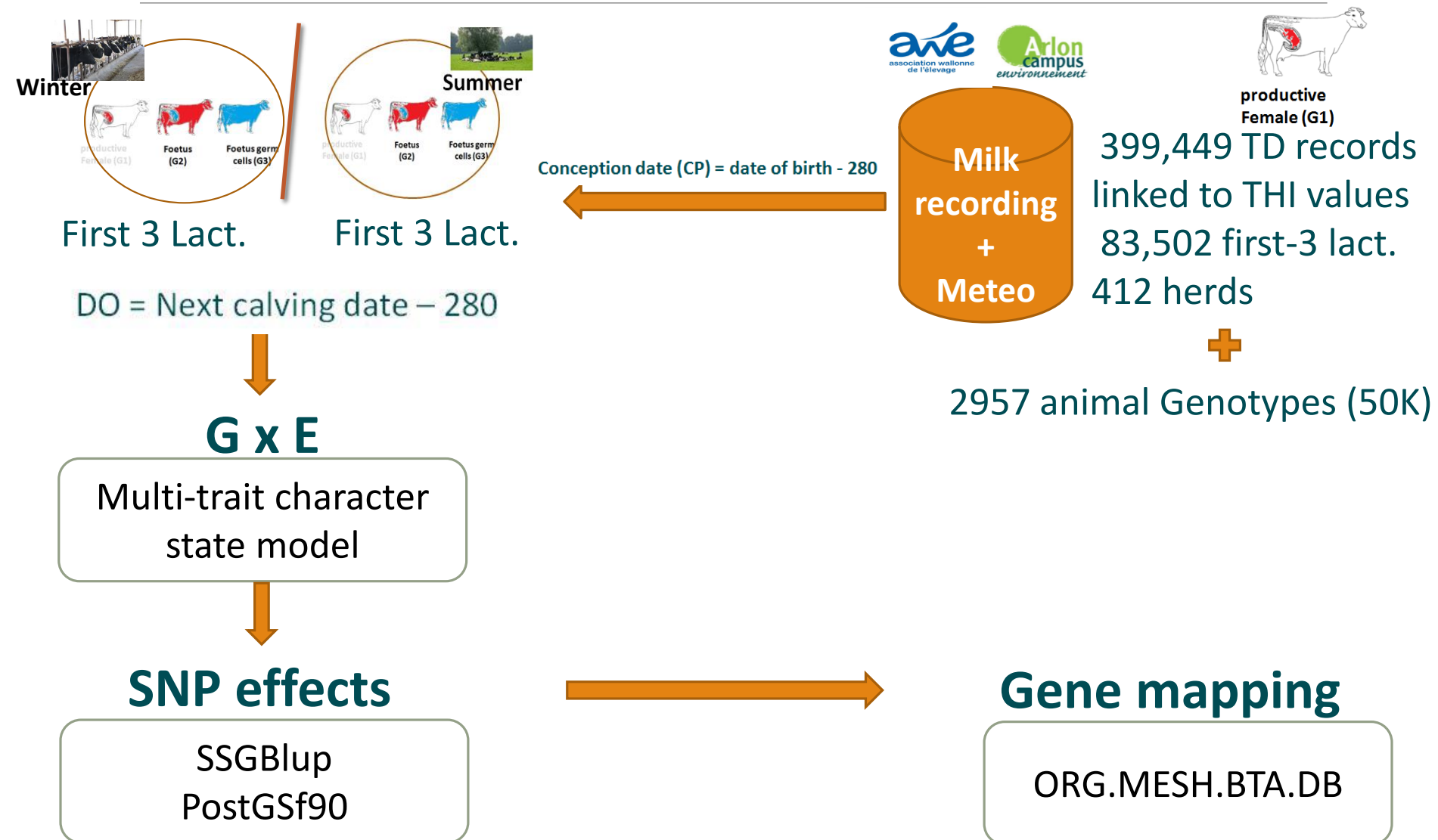
# Objectives

- **G x HS exposure** during the fetal development period



- Characterize **genomic regions** responsible for HS exposure at early life as affecting lifetime fertility

# Data & models



# Genetic correlations



Winter

Summer

	Winter			Summer		
	L1	L2	L3	L1	L2	L3
L1	.07	.77	.71	.73	.61	.62
L2		.06	.74	.75	.68	.60
L3			.06	.70	.58	.58
L1				.07	.68	.62
L2					.05	.63
L3						.05



Winter



Summer



Winter

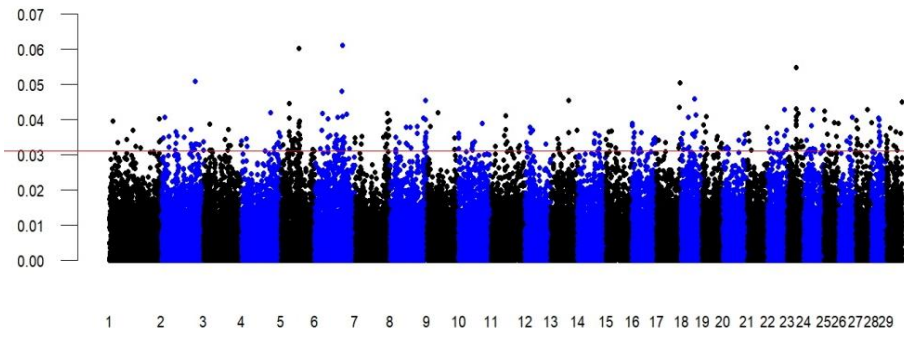
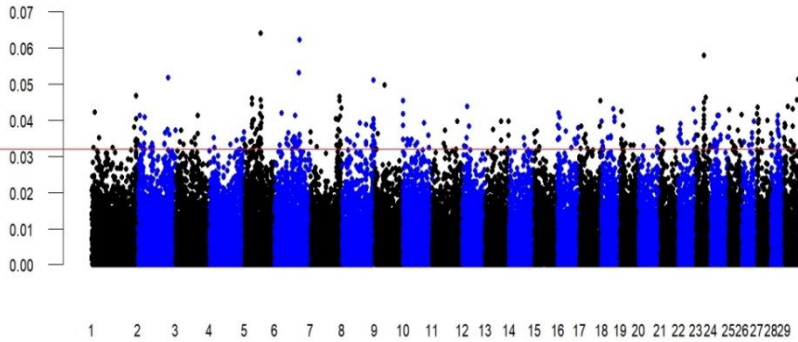


# Manhattan plots



Summer

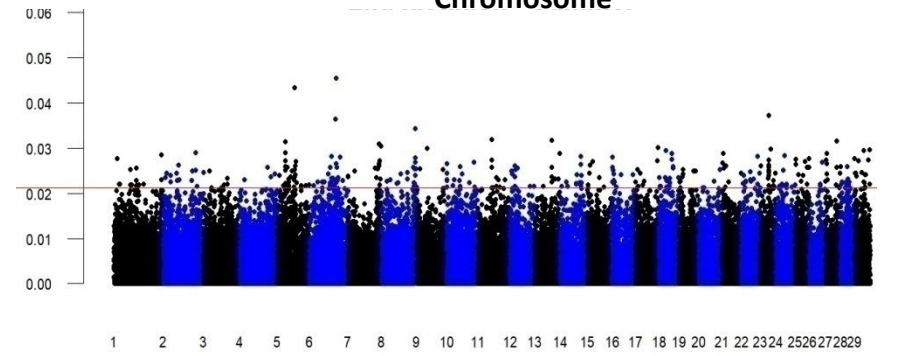
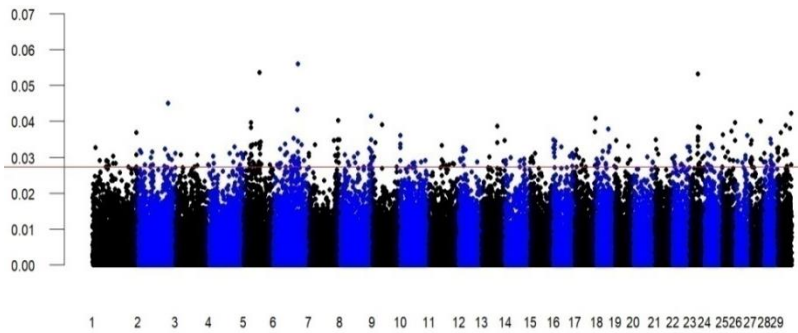
L1



Chromosome

Chromosome

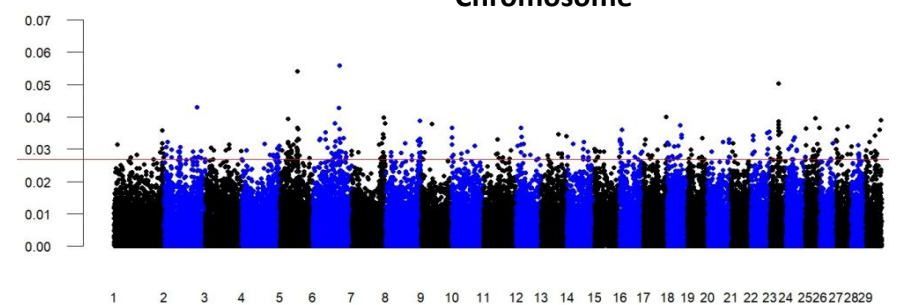
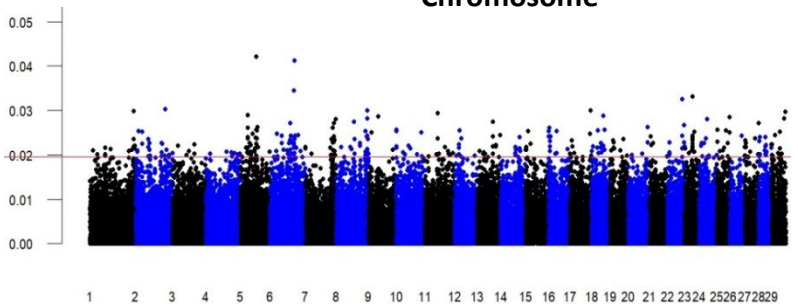
L2



Chromosome

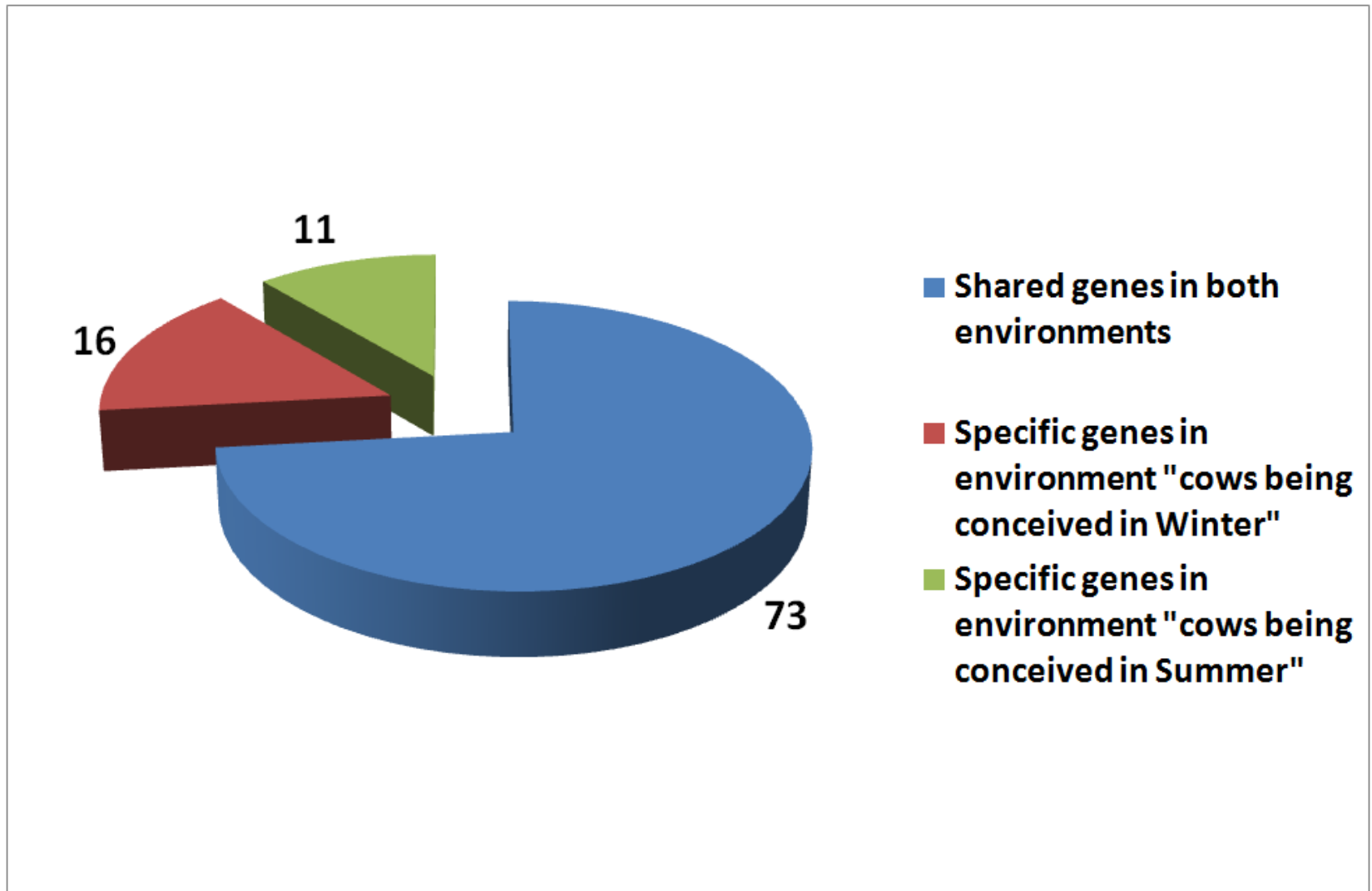
Chromosome

L3





# Proportions of identified genes



# Functional gene enrichment (shared)

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<b>MeSH ID</b>	<b>Term</b>	<b>Number of signif. genes</b>	<b>Total genes</b>	<b>P-Value</b>
<b>D011270</b>	Pregnancy, Animal	12	71	0.001
<b>D013312</b>	Stress, Physiological	6	23	0.002
<b>D030762</b>	Estrous Cycle	12	86	0.004
<b>D047109</b>	Fetal Development	3	7	0.006
<b>D044127</b>	Epigenesis, Genetic	6	29	0.006

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# Functional gene enrichment (specific summer)



Summer

MeSH ID	MeSH term Name	Signif. Genes	P-value
D050883	HSC70 Heat-Shock Proteins	4	0.003
D019791	Guanidine	4	0.006
D018384	Oxidative Stress	9	0.009
D007335	Insulin-Like Growth Factor II	4	0.014

# Functional gene enrichment (specific winter)



Winter

MeSH ID	MeSH term Name	Signif. Genes	P-value
D001120	Arginine	10	0.001
D051766	Early Growth Response Protein 1	3	0.017
D004847	Epithelial Cells	19	0.025
D000222	Adaptation, Physiological	3	0.029

# Take home messages

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- $r_g$  correlations across both early-life environments were lower than .80
- Differentially expressed genes when offspring being conceived in summer compared to winter
- Ongoing work in the identification of potential gene candidates for metabolism and physiology traits

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