

Quantitative trait loci (QTL) for body weight in Arctic Charr reared in different environmental conditions



Aquaculture in Canada

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- ❑ Dominated by Atlantic salmon (*Salmo salar*)
- ❑ Decreased market value
- ❑ Overproduction and competition
- ❑ Alternative species



Charr Projects in Atlantic Canada

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- Overall goal = commercial broodstock
 - ▣ Salinity tolerance
 - ▣ Flesh pigmentation
 - ▣ Cryopreservation
 - ▣ Sex control
 - ▣ Triploid induction
 - ▣ **Genomic evaluation**



Arctic Charr Aquaculture

□ Benefits:

- High stocking density
- Grows well in cold water
- “luxury” food item

□ Problems:

- Variable growth rates
- Early maturation
- Decreased flesh quality



How can Genomics help?



- Genetic evaluation of a broodstock can help identify individuals who are predisposed to a trait of interest
- When genetic selection is combined with traditional selection, selection intensity may increase rapidly

Analysis of a quantitative trait (QTL)

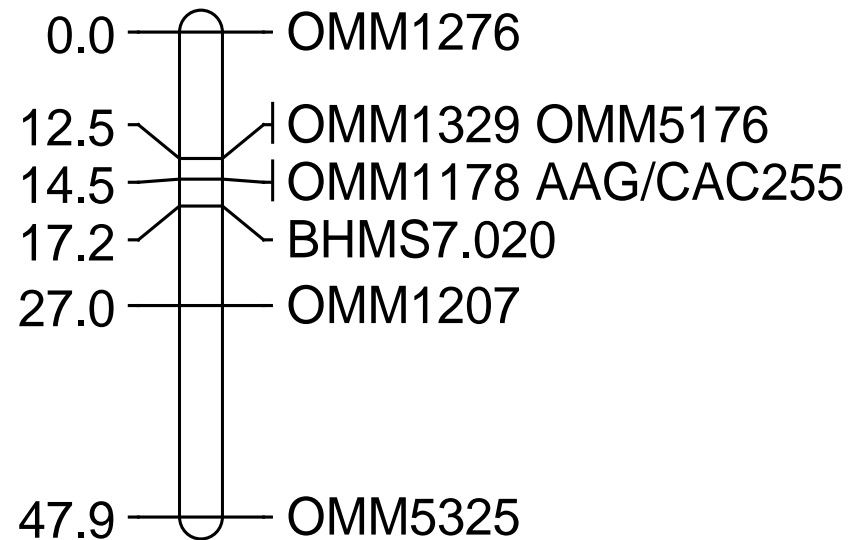


- QTL- genomic regions containing alleles that affect a quantitative trait (ie: body weight)
- If a marker is linked to a region coding for trait of interest – there is a statistical association between the phenotype and the alleles

Analysis of a quantitative trait (QTL)

AC-32f

- Microsatellite markers – sequence repeats with variation in number of repeats
- Markers selected from QTL studies on AC, AS & RT

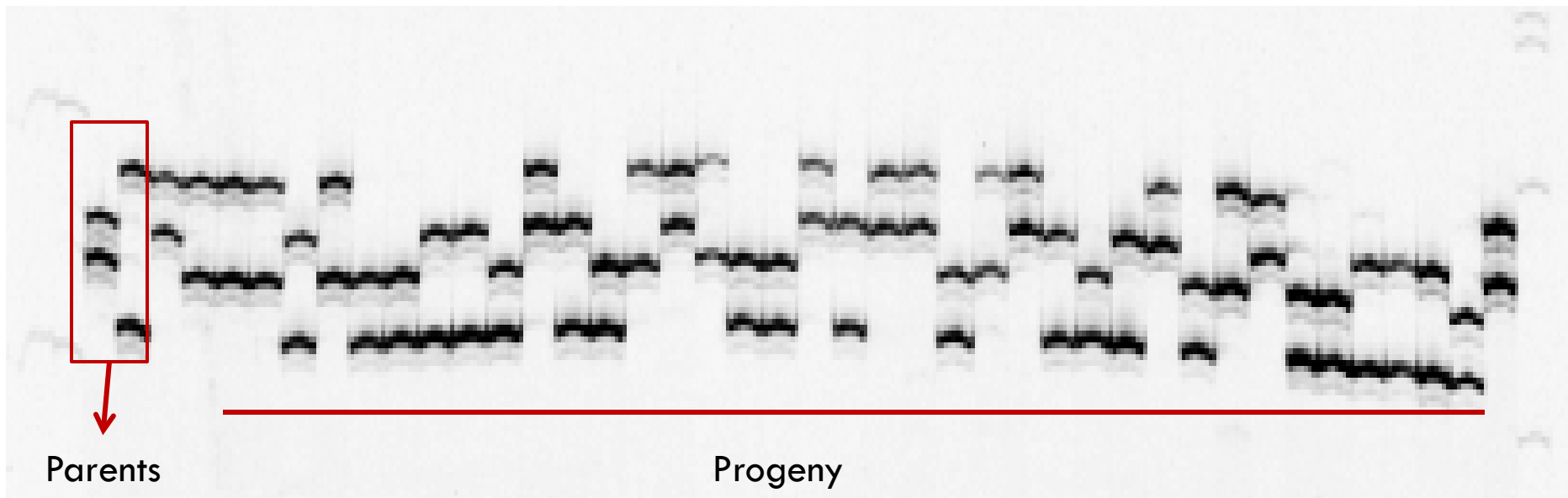


Analysis of quantitative trait (QTL)



Analysis of a quantitative trait (QTL)

- Collect tissue samples, extract DNA
- Measure the trait of interest
- PCR reaction with marker-specific primers
- Visualize on acrylamide gel

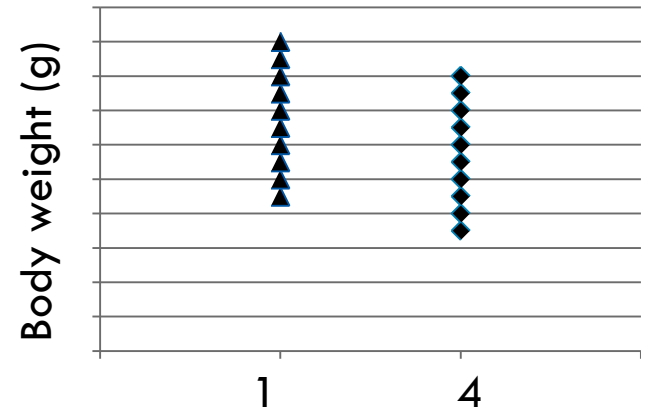
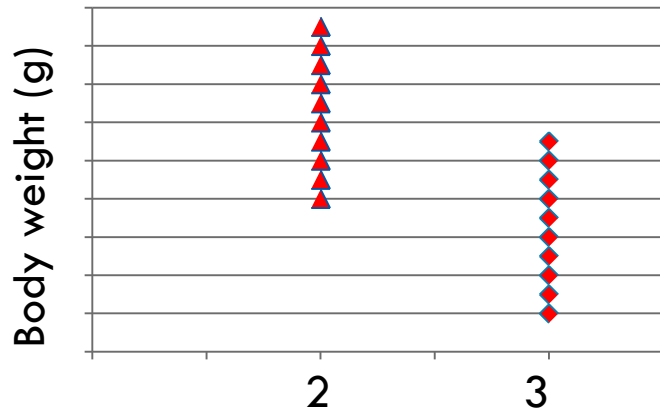
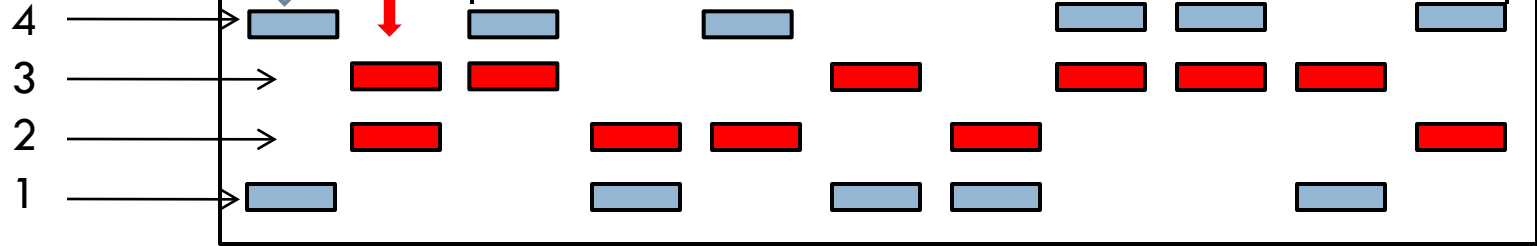


Analysis of quantitative trait (QTL)



progeny

alleles

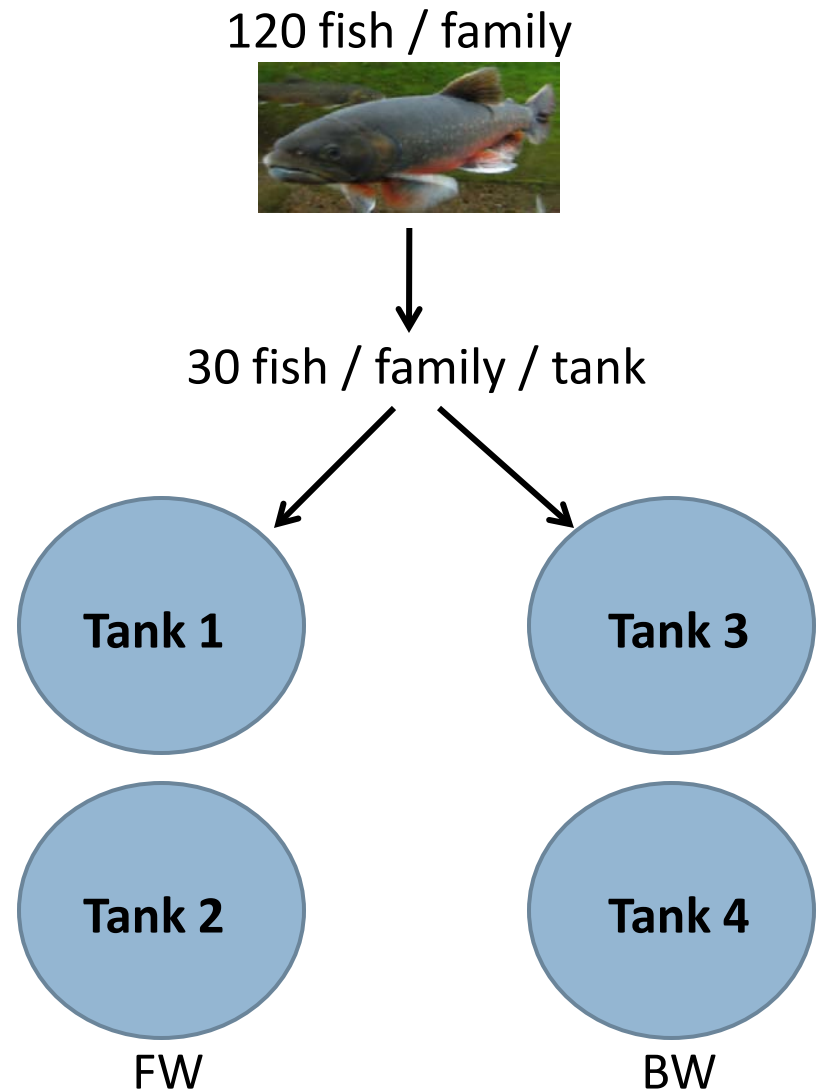


Background

- Pilot study found significant QTL for body weight on AC-8 (Moghadam et al. 2007)
- This study investigated 2 experimental Fraser strain families
- We wanted to see if this result would also be found in a larger number of families grown at a commercial site

Methods

- 30 full-sib families
- PIT-tag 120 fish from each family (n=3600)
- 4 - 16 m³ tanks
- 2 environments (FW & 20 ppt BW)



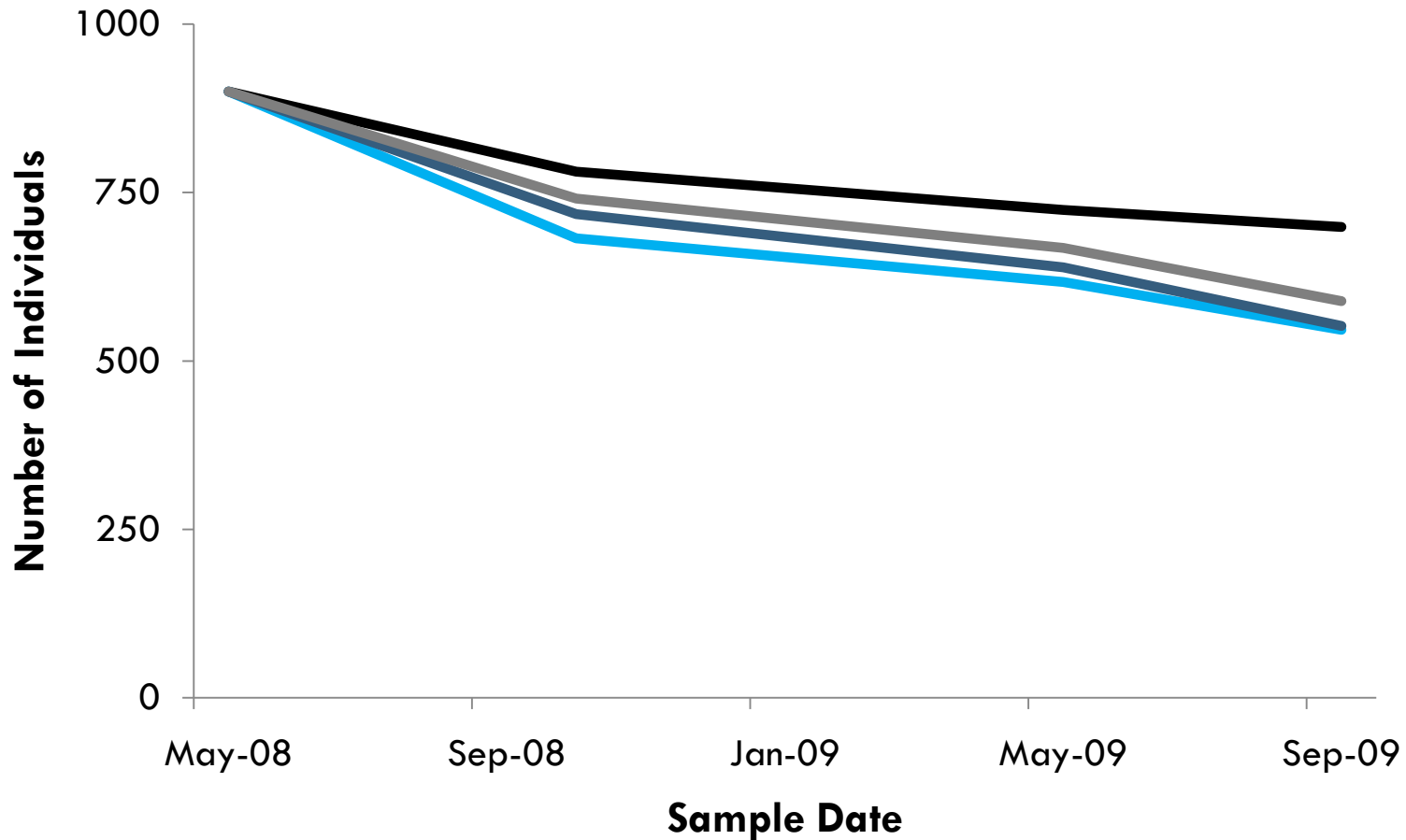
Can Aqua – Advocate Harbour



Progress

- 3600 fish were PIT-tagged, weighed, measured and transferred to CanAqua in February 2008
- Tissue samples collected
- All individuals genotyped for QTL markers
- Measured body weight, length and survival at 5 sampling dates

Preliminary Results - Survival



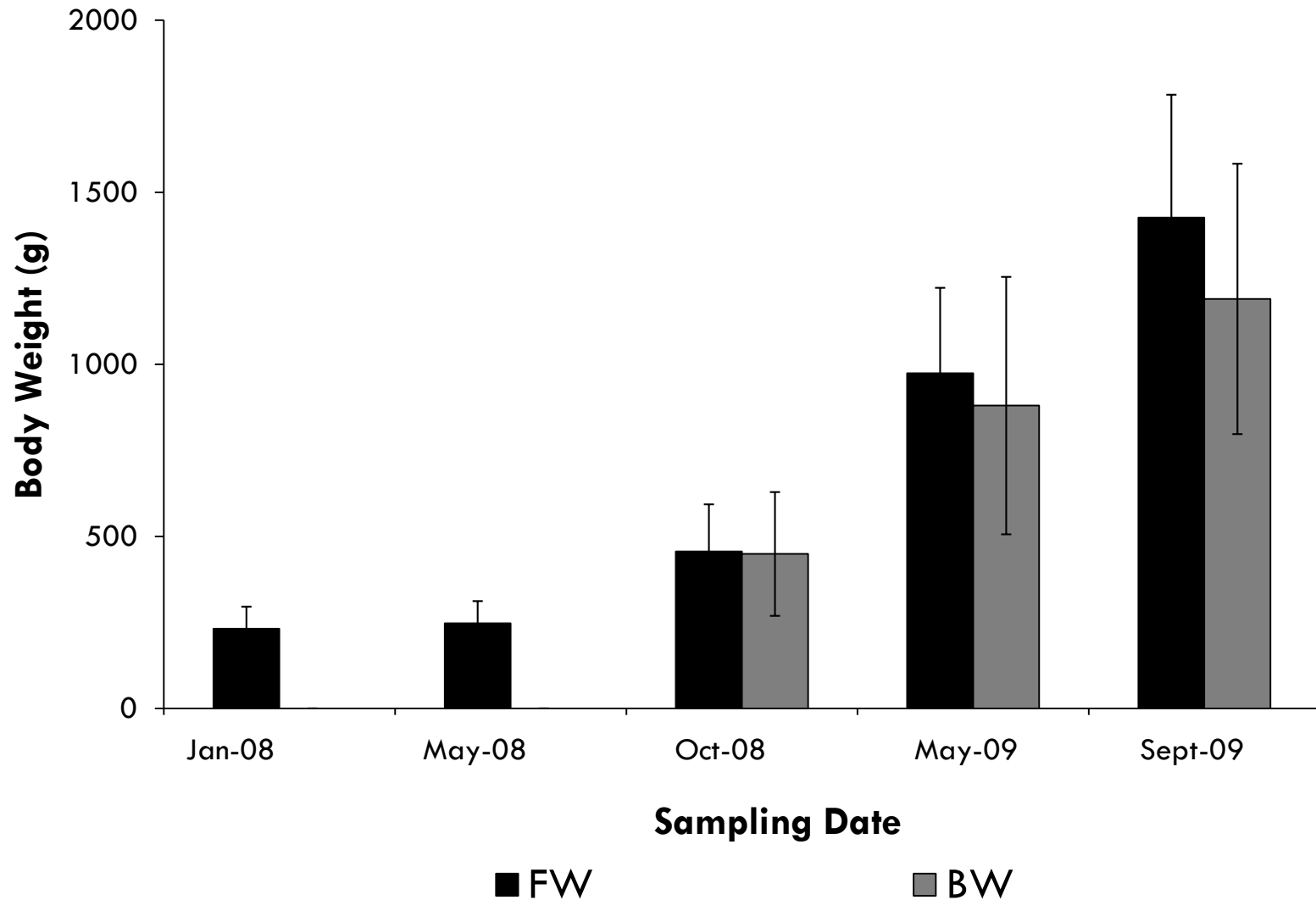
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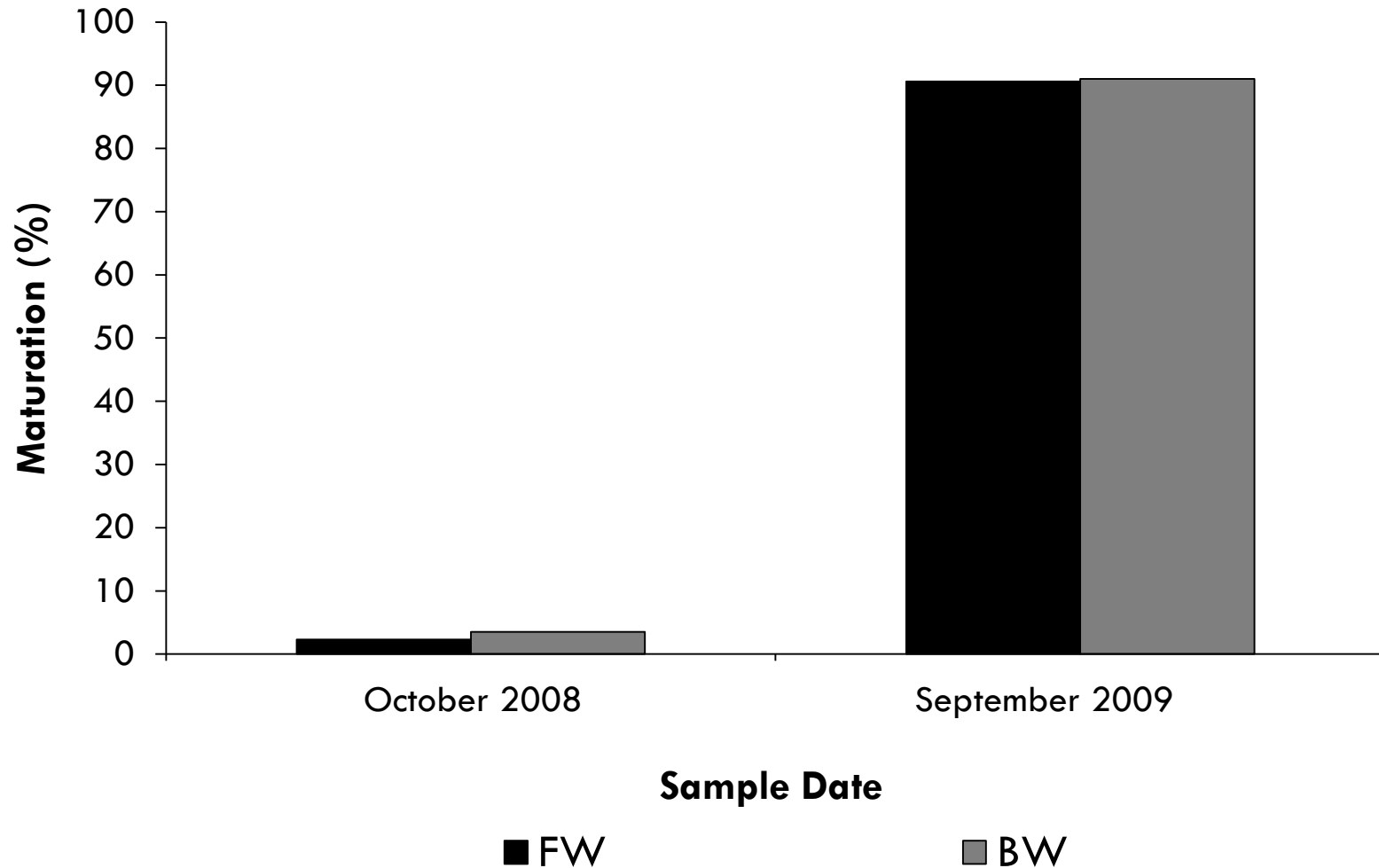
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Preliminary Results – Body Weight



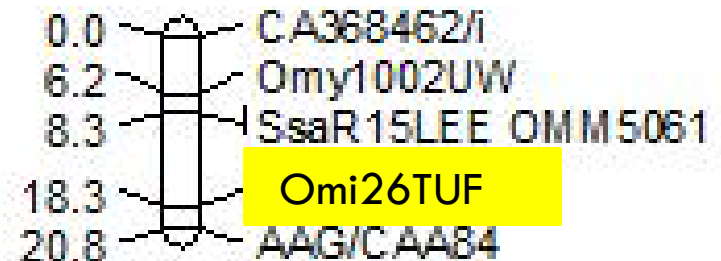
Preliminary Results - Maturation



Preliminary Results - Genetics

- Most significant QTL is on AC-8
- Found in multiple families in both males and females
- QTL accounts for 8-17% of the variation in body weight

AC8



Application & Goals

- Significant QTLs can help direct selection through Marker Assisted Selection (MAS)
- Overall goal = select for fast growth & reduced early maturation
- The specific objectives of this project are to determine if QTL for body size and early maturation already identified from genome scans have detectable effects across families and environments

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