Fitness Studies – SEAK Chum Salmon Progress on genetic markers



Kyle Shedd, E. Lescak, H. Hoyt, C. Habicht
Alaska Department of Fish and Game Gene Conservation Lab
Alaska Hatchery Research Program Informational Meeting
March 7, 2019

Alaska Hatchery Research Program

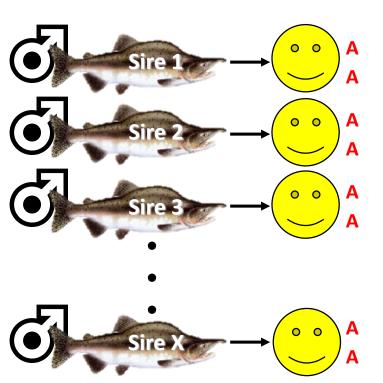
- 1) What is the genetic structure of pink and chum in PWS and SEAK?
- 2) What is the extent and annual variability of straying?
- 3) What is the impact on <u>fitness</u> (productivity) of natural pink and chum stocks due to straying hatchery pink and chum salmon?

Genetic marker needs

- Similar needs to pink salmon markers
- ~300 genetic markers
- High information content
 - > 0.3 average minor allele frequency (MAF)

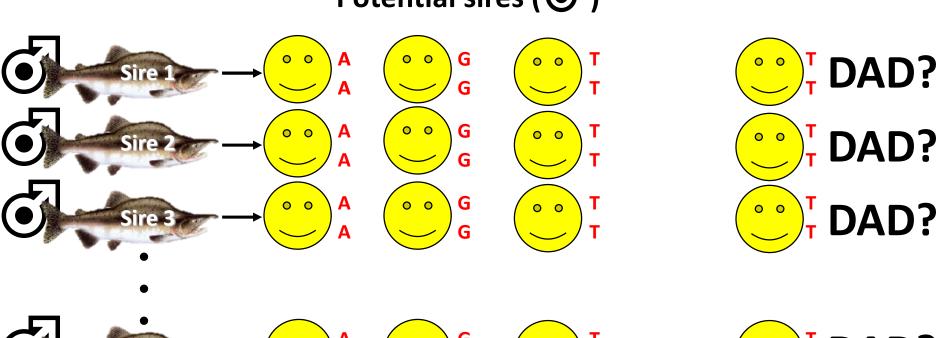
Markers



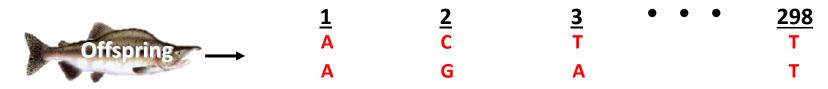


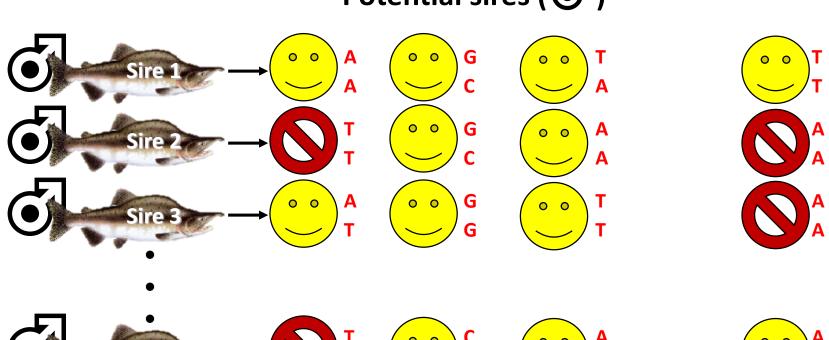
Markers





Markers

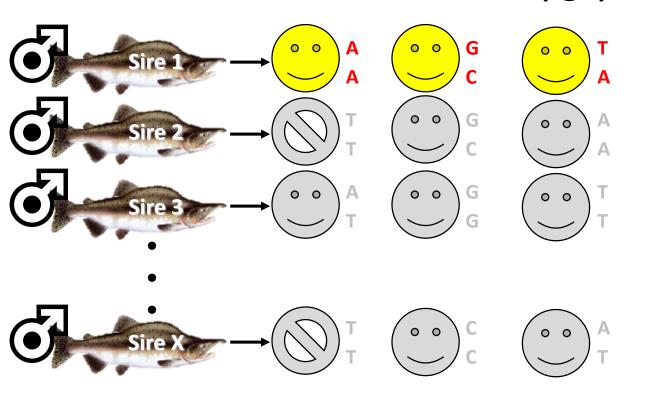




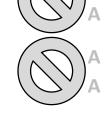


Markers









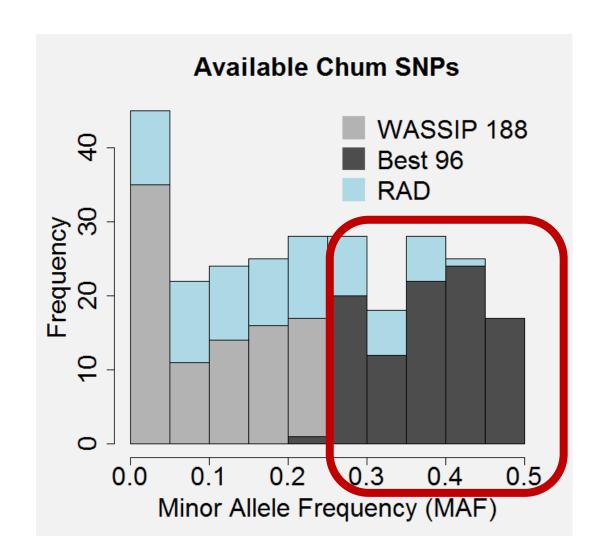


Genetic marker needs

- Similar to pink salmon
- ~300 genetic markers
- High information content
 - > 0.3 average minor allele frequency (MAF)
- New, high throughput chemistry
 - GTseq panel

Genetic markers currently available

- Old chemistry
 - "WASSIP 188"
 - "RAD 72"
 - UW



Future work

- Current markers are not adequate
- Find or develop chum GTseq panel
 - ~300 markers
 - MAF > 0.3
- Harvest existing RAD data from Western Alaska project
- Genetics technology changes
- Waiting saves \$\$\$

