Chum Salmon progress on genetic markers



Kyle Shedd Gene Conservation Laboratory Alaska Department of Fish and Game AHRP Informational Meeting March 6, 2020

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)

Genetic markers for parentage analysis

Markers

<u>1</u> A

Α



<u>2</u>	<u>3</u>	• • •	<u>298</u>
G	т		т
G	Т		т

Potential sires (







Genetic markers for parentage analysis Markers <u>2</u> C 298 T <u>3</u> T <u>1</u> A G Potential sires () G 0 0 Α 0 0 0 0 Δ 0 0 G T 0 0 Α G 0 0 0 0 0 0 т G 0 0 T С 0 0



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 GT-seq panel
 - ~300 markers
 - MAF > 0.3
- Try 350 SNP GT-seq panel developed in Washington
 - Send SEAK samples, find out MAF
- Genetics technology changes
- Waiting saves \$\$\$

Questions?