Genetic Stock Composition of the Commercial Harvest of Chinook Salmon in the Copper River District, 2013-2017

A Report to the Alaska Board of Fisheries

December 2017



Sara Gilk-Baumer Commercial Fisheries Division Gene Conservation Laboratory

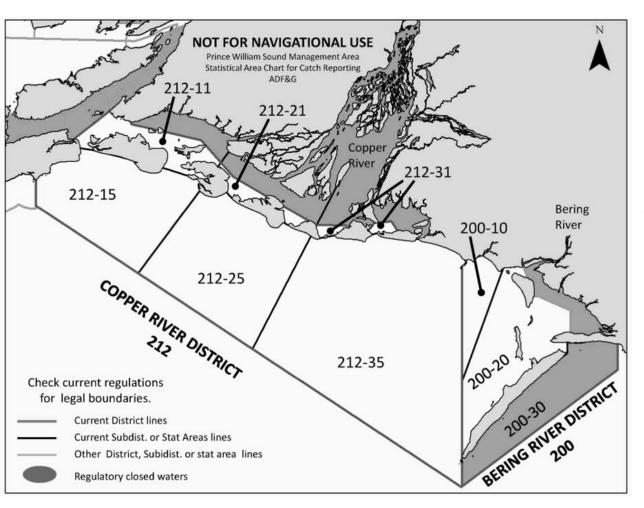
Outline

- Background
- Sampling methods
- Mixed stock analysis methods
- Results
- Summary and conclusions

Background

- Chinook Salmon Research Initiative (CSRI)
 - Copper as 1 of 12 indicator stocks
 - Information gaps identified
 - Adult spawning abundance
 - Juvenile abundance
 - Stock of origin of marine catches
- Project proposed for 2013–2017 to identify stock of origin in marine commercial fisheries in Copper River District

Background: Copper River District



- Harvest of local and non-local stocks
- Management strategy: provide inriver passage from all stocks

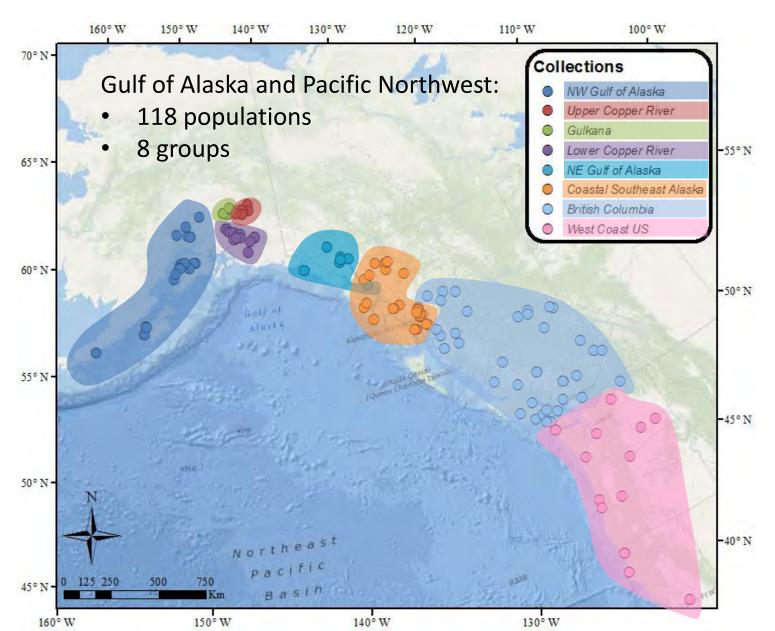
Sampling Methods

- Chinook salmon from commercial gillnet harvest in marine waters
- By period, 5-6 weeks beginning mid-May
- Sample size: 200 samples per period
- Genetic tissue: axillary process
- Some paired with age (scale), sex, and length





Mixed Stock Analysis: Genetic Baseline

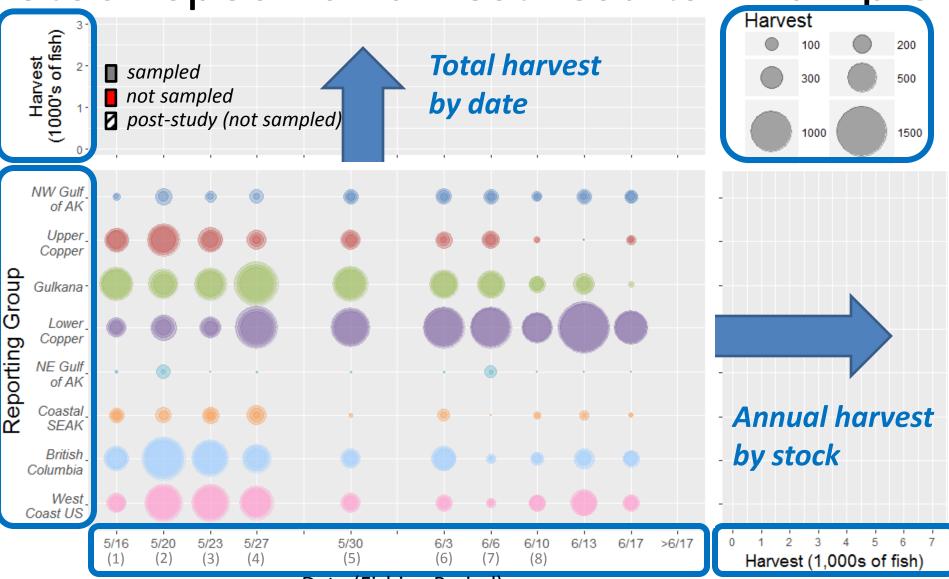


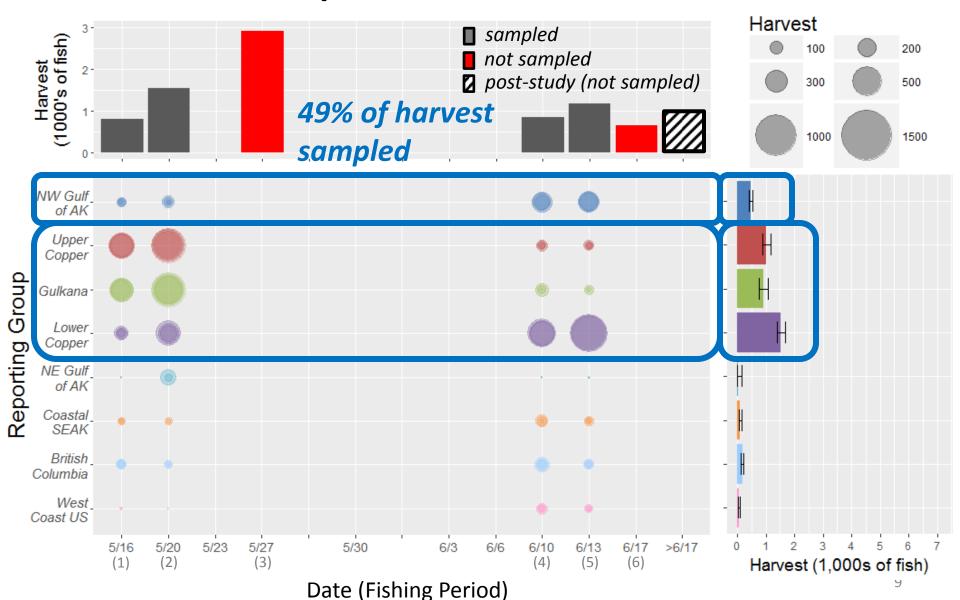
Mixed Stock Analysis Methods

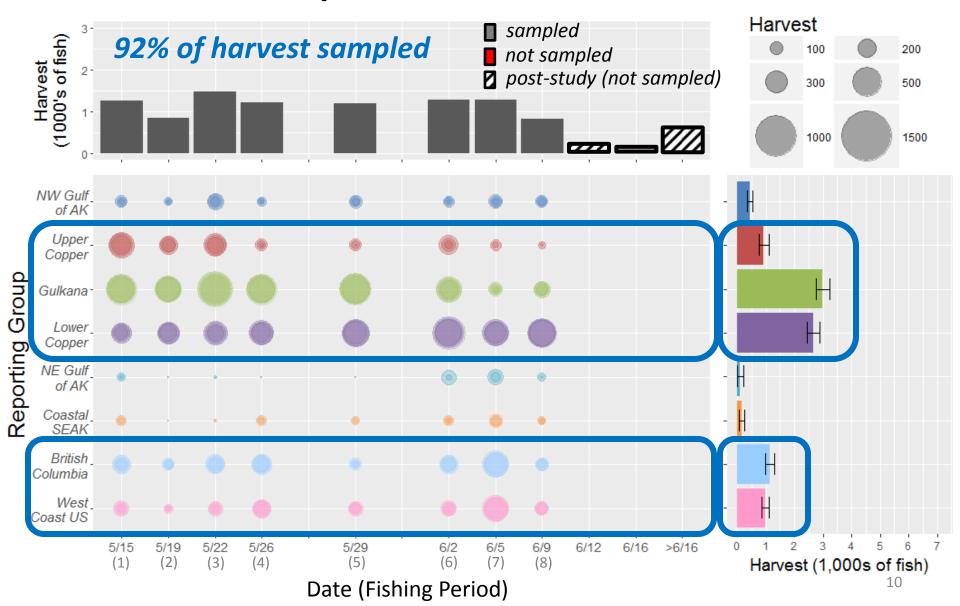
- Harvest samples genotyped for 43 SNP markers in common with baseline
- Estimate relative stock composition

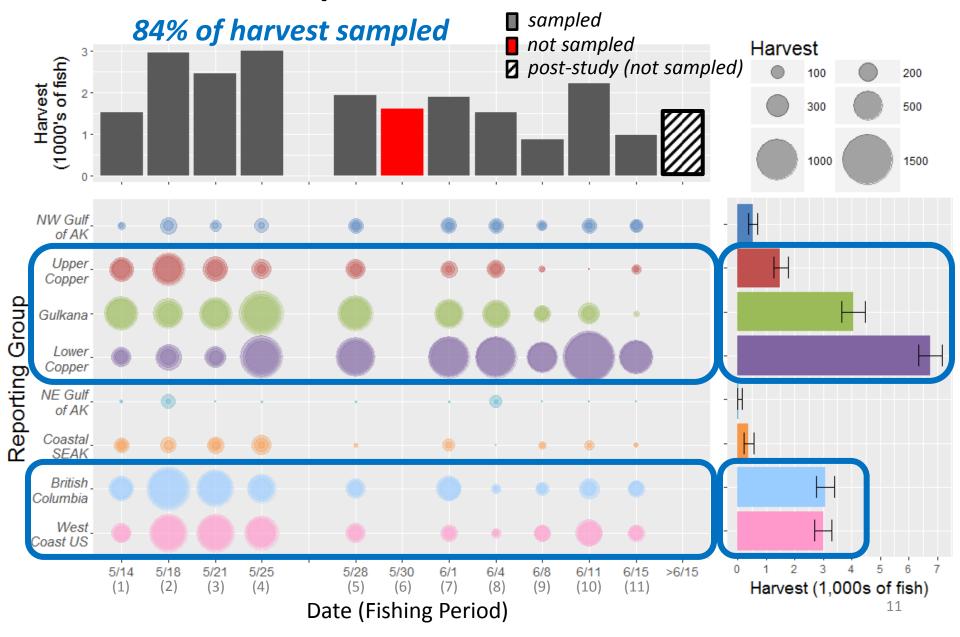


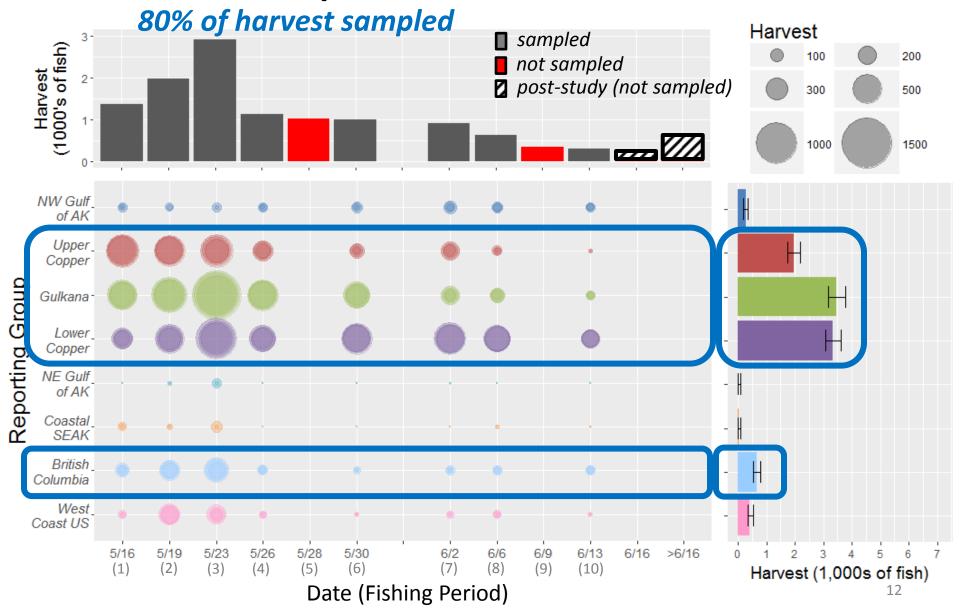
Stock-specific harvest results: Example

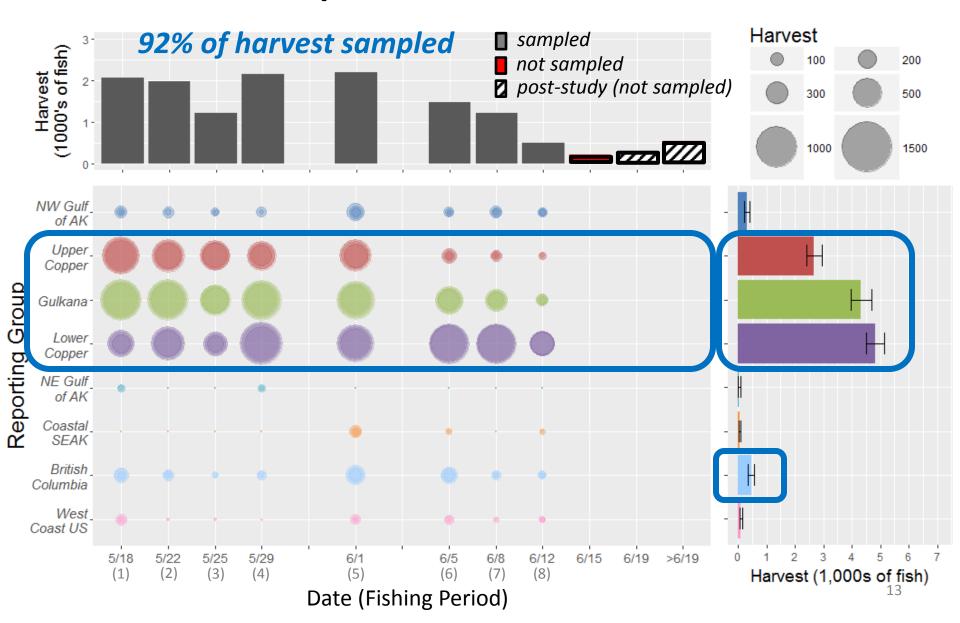




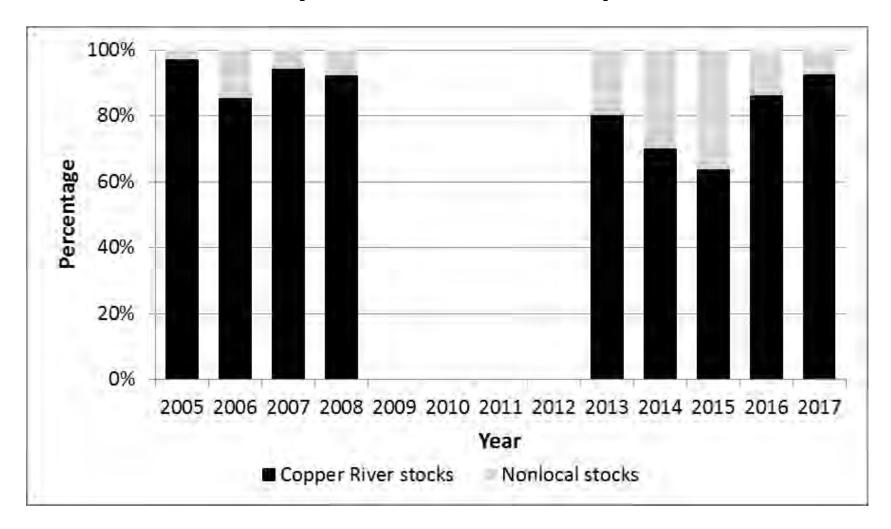






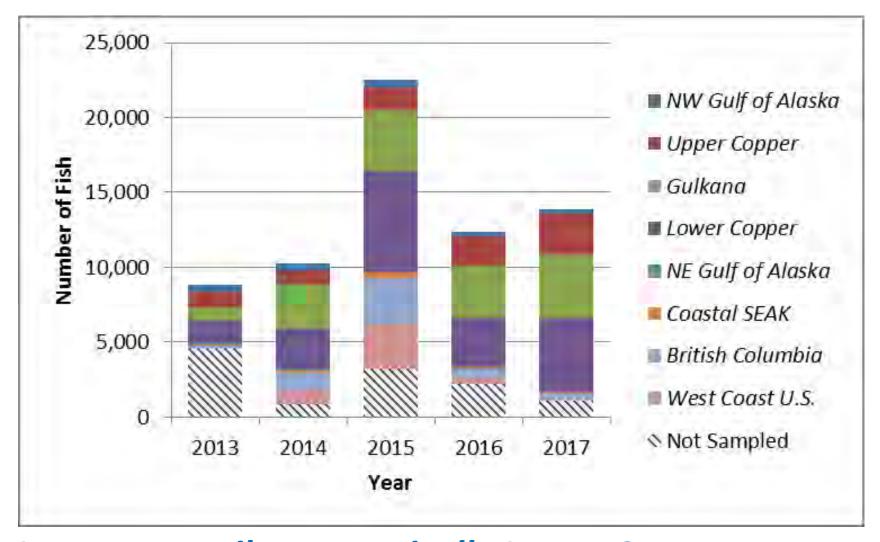


Summary: Stock Composition



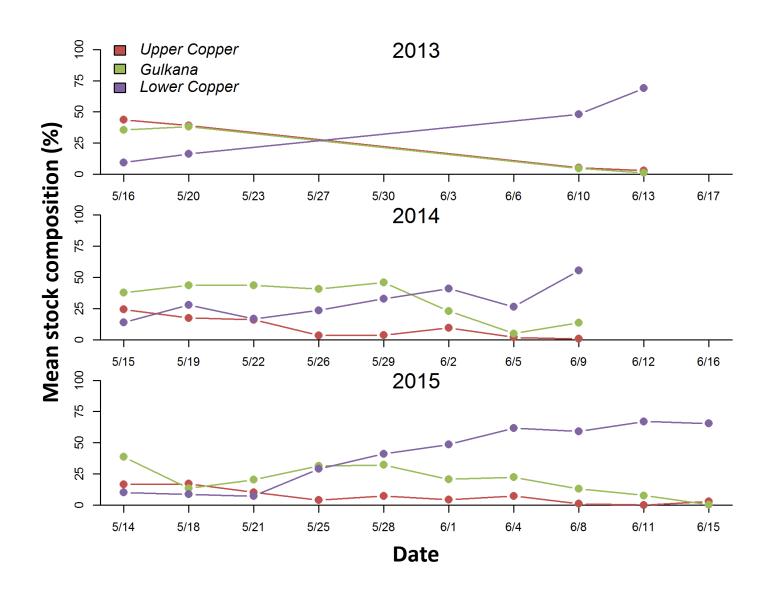
Fishery dominated by Copper River stocks

Summary: Stock Composition

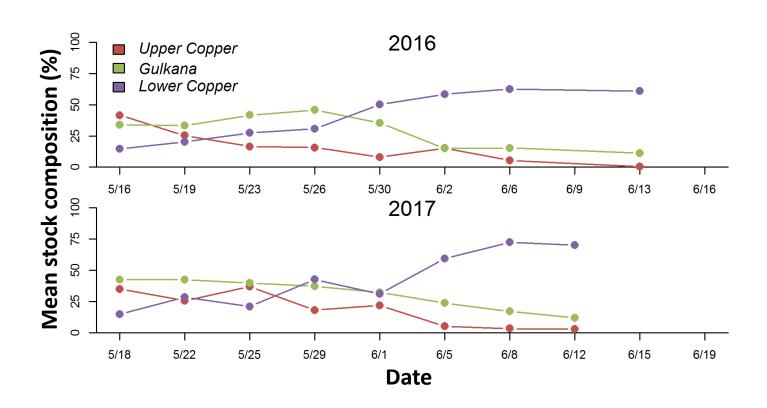


Largest contributor: typically Lower Copper group 15

Summary: Run Timing



Summary: Run Timing



Upper Copper group declined,

Lower Copper group increased

Conclusions

- Caveats:
 - Small contributions with large relative error
 - Not all strata sampled
- Run timing patterns useful for management
- Most harvest originated from populations in the Copper River
- Estimation of non-local stocks could provide information for production models

Acknowledgements

- Chinook Salmon Research Initiative
- Co-Authors
 - Dani Evenson, Kyle Shedd, Heather Hoyt, Chris Habicht, Bill Templin, Stormy Haught
- GCL staff
 - Paul Kuriscak, Erica Chenoweth, Zac Grauvogel, Chase Jalbert, and Zach Pechacek
- Cordova staff
 - Jeremy Botz, Stacy Vega, Steve Moffitt, Jim O'Rourke,
 Angie Kelly, and Scott Prevatte

