

NOAA FISHERIES



College of Fisheries and Ocean Sciences



Updating bycatch genetics workflow

Patrick Barry
University of Alaska
College of Fisheries and Ocean Science

Who I am

- UAF CFOS
 - PhD. work with Tony Gharrett and Megan McPhee.
 - Population genetics of sockeye salmon in SE Alaska.

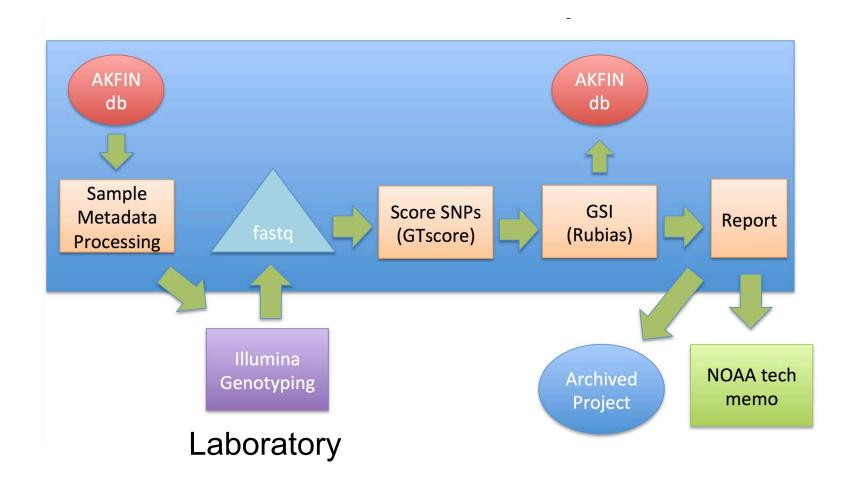


• Skillset:

 Population genetics, quantitative genetics, eDNA, R coding, molecular biology



Data Analysis Pipeline





AKFIN database

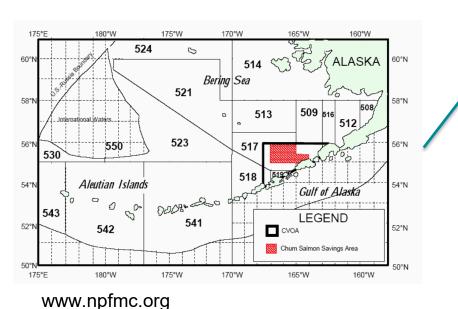
Alaska Fisheries Information Network
Established under direction of Pacific
States Marine Fisheries Commission (PSMFC)
Goal:

- Link observer records to genetic and age information (confidentially)
- Facilitate data exploration and spatio-temporal modeling (PCCRC funded project).



AKFIN database: Trivial example

 Do static closures preserve the same stocks of fish from year to year?





- Pull genetic records
- Format mixture
- MSA rubias

Spatio-temporal dynamics of chum salmon PSC

- Quantify the inter- and intra-annual patterns of genetic stock composition across space, time, fishing sector, and fish age
- Evaluate how age- and stock-specific distributions of chum salmon vary with biotic and abiotic factors in the Bering Sea.
- End goal: Move to predictive framework for bycatch avoidance



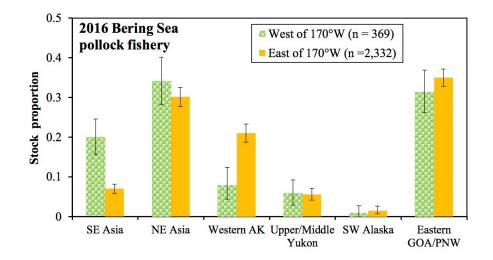
Data visualization: R shiny (interactive web) application

pdbarry.shinyapps.io/gsi_salmon_comparisons/



Genetic Stock Composition Analysis of Chum Salmon from the Prohibited Species Catch of the 2016 Bering Sea Walleye Pollock Trawl Fishery and Gulf of Alaska Groundfish Fisheries

J. A. Whittle, C. M. Kondzela, Hv. T. Nguyen, K. Hauch, D. Cuadra, and J. R. Guyon

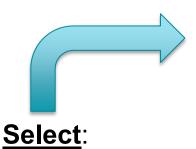


U.S. DEPARTMENT OF COMMERCE
National Oceanic and Atmospheric Administration
National Marine Fisheries Service
Alaska Fisheries Science Center

January 2018



Data visualization



Region

- Bering Sea / GOA
Spatial strata

Spatial strata

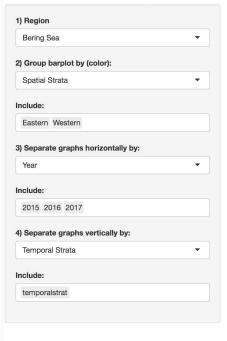
- NMFS area
- East/West 170

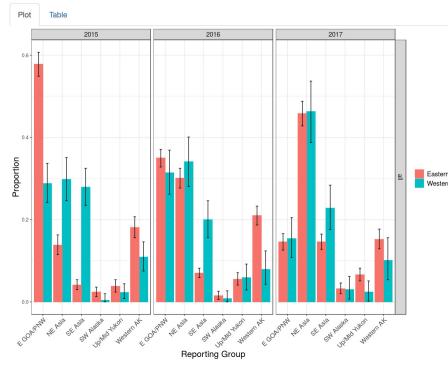
Temporal strata

- Early/Middle/Late
- A/B season

Year

Genetic Stock Composition of salmon bycatch



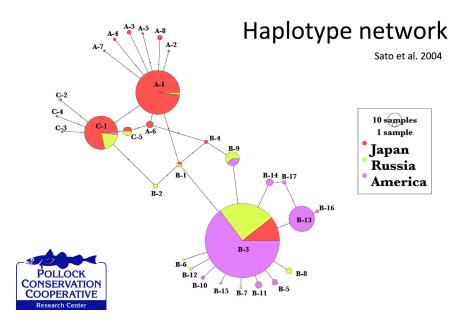


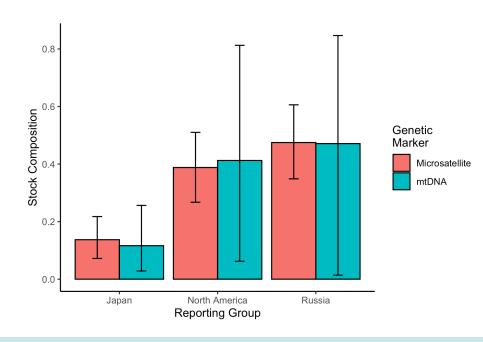


Shipside stock identification of chum salmon bycatch with nanopore sequencing

 Can we estimate Asia vs. North America contributions to mixtures?









Acknowledgements

NOAA ABL

Wes Larson, Chris Kondzela, Chuck Guthrie, Jackie Whittle, Katie Damelio, Jordan Watson, & Ellen Yasumiishi

AKFIN

Camille Kohler Rob Ames Robert Ryznar

UAF

Megan McPhee

