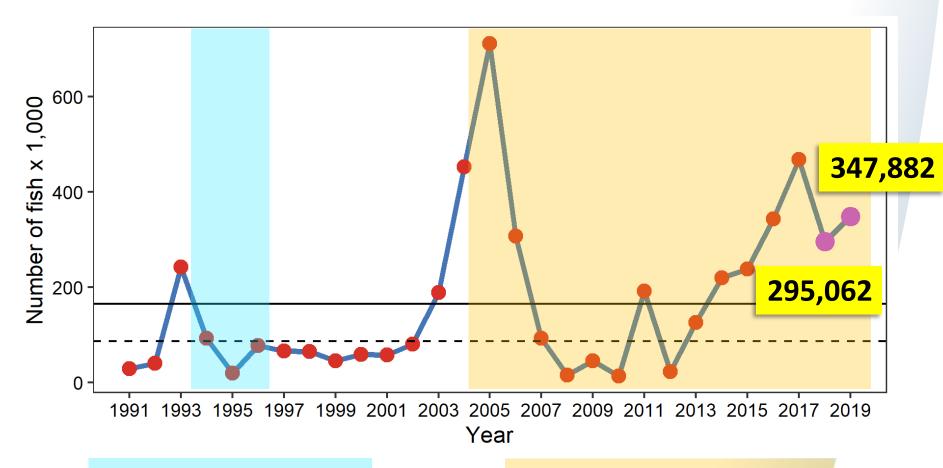


Genetic stock composition analysis of chum salmon from the prohibited species catch of the 2018-2019 Bering Sea walleye pollock trawl fishery and Gulf of Alaska groundfish fisheries

C. Kondzela, P. Barry, J. Whittle, J. Watson, E. Yasumiishi, and W. Larson

Auke Bay Laboratories Juneau, Alaska

NPFMC meeting April 2021 Virtual

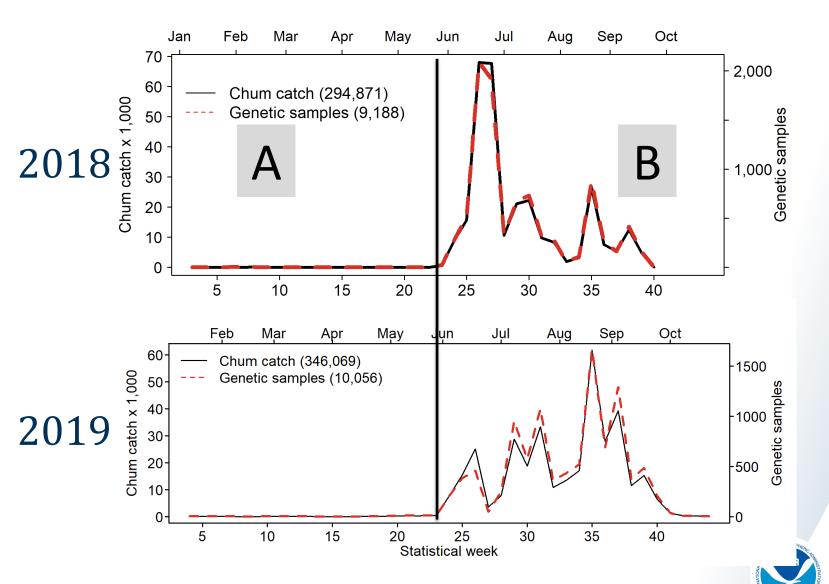


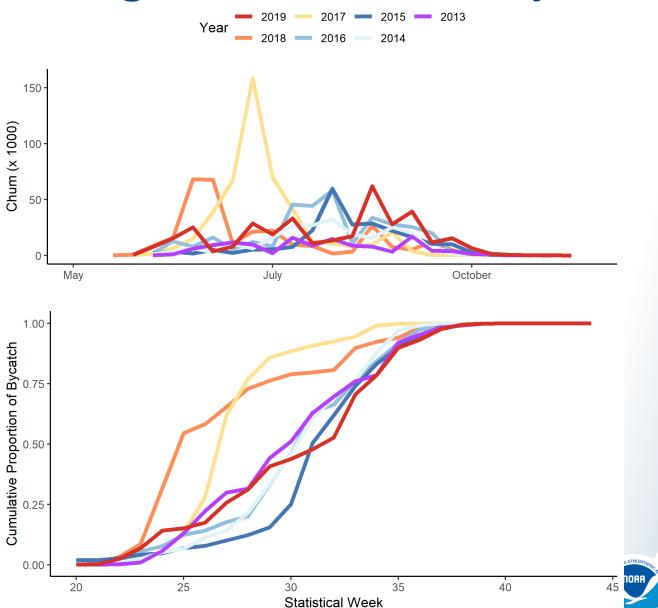
1994: Scales

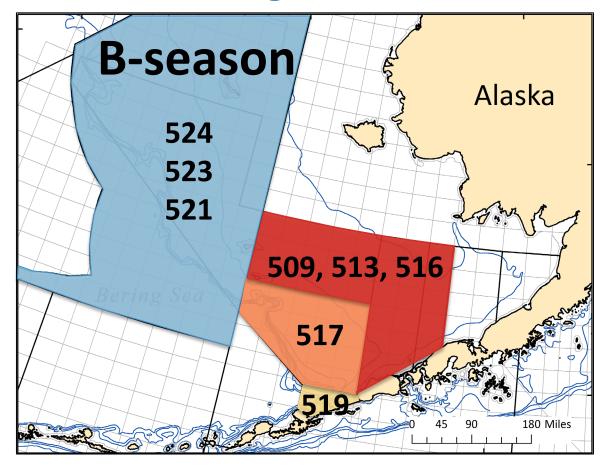
1994 – 1996: Allozymes

2005 - 2019: Microsatellites

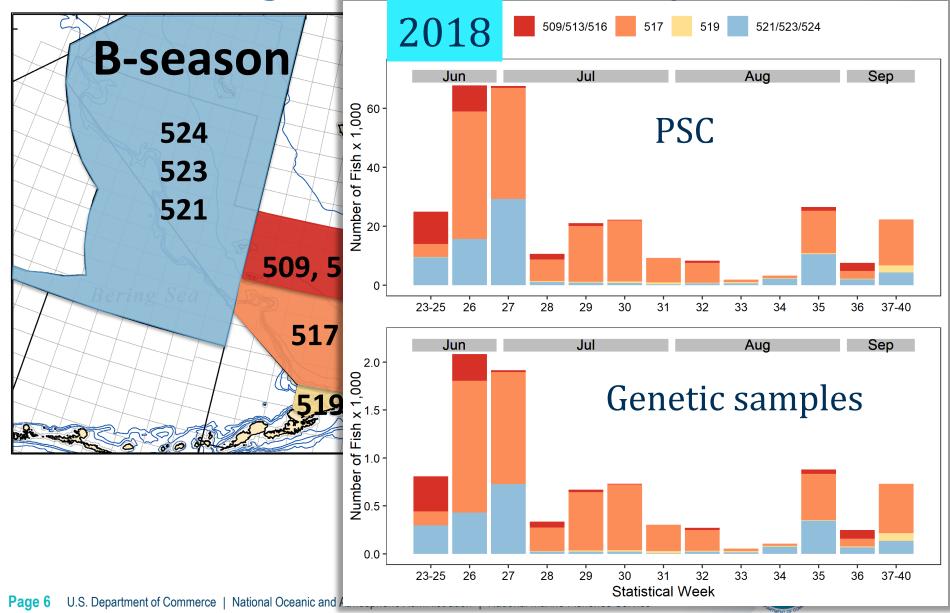


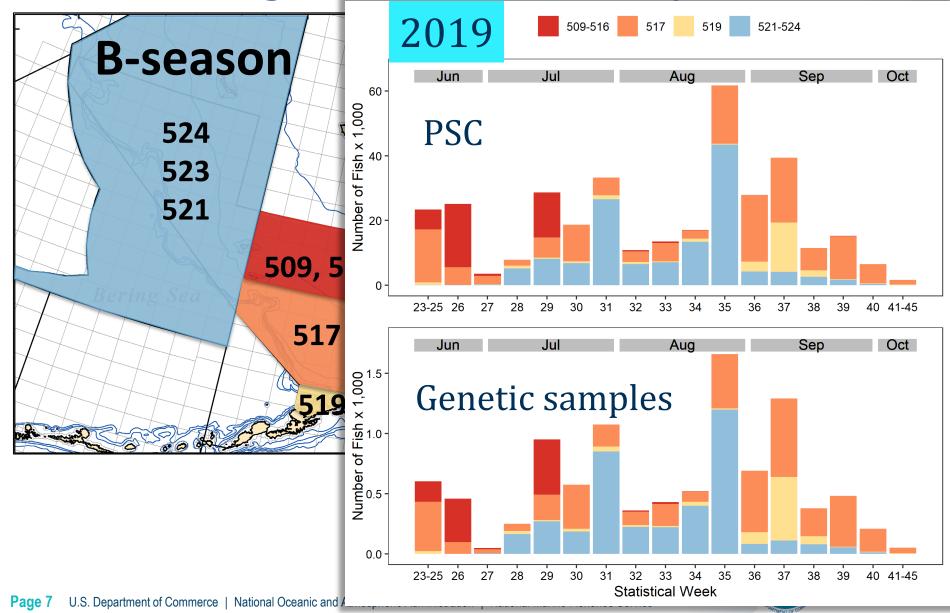




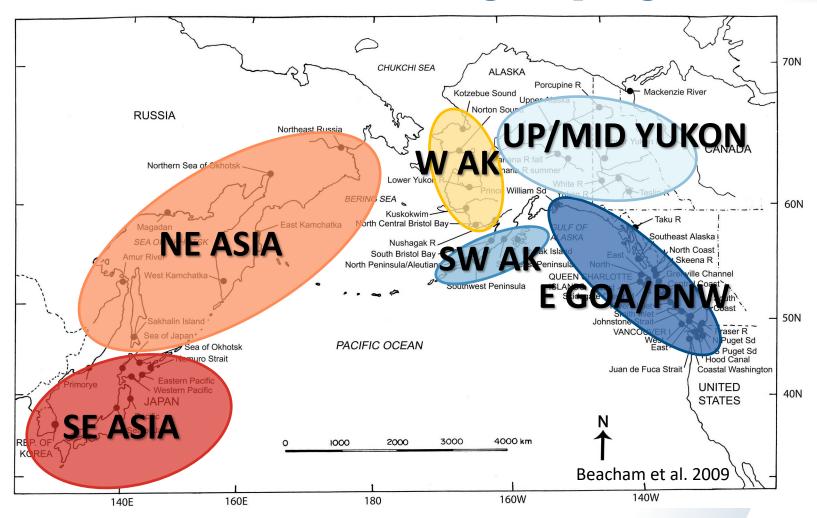




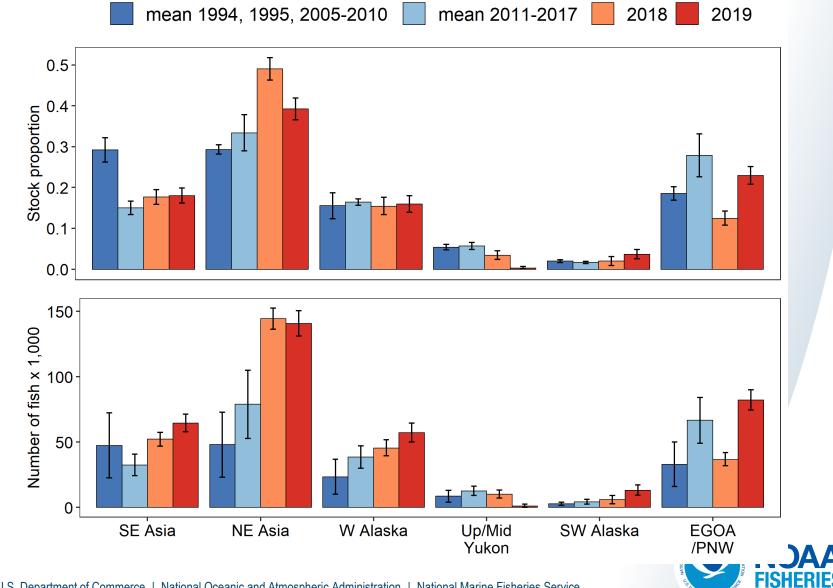


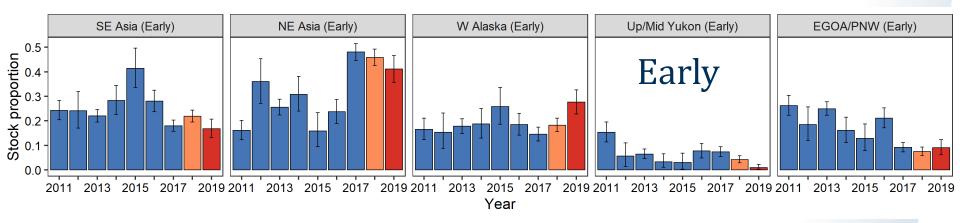


Chum salmon stock groupings

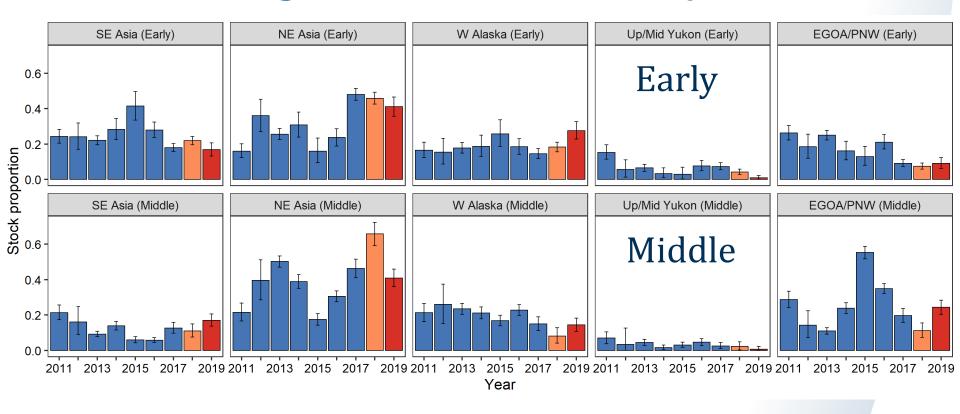




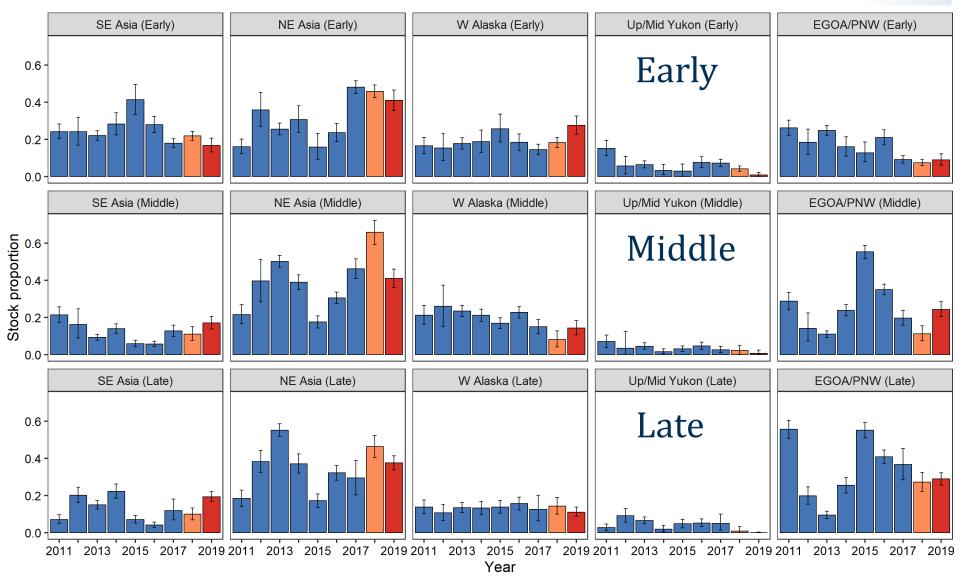






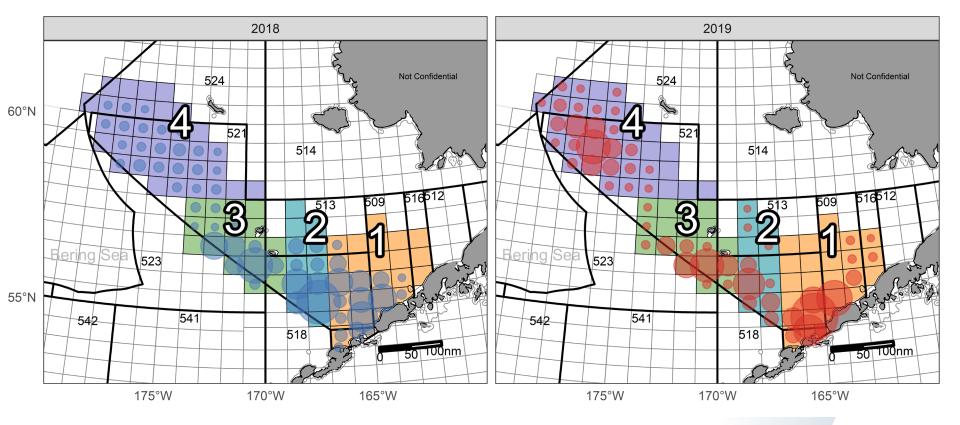




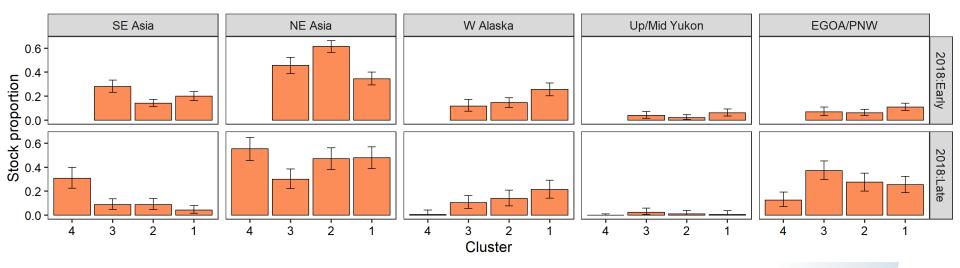


2018

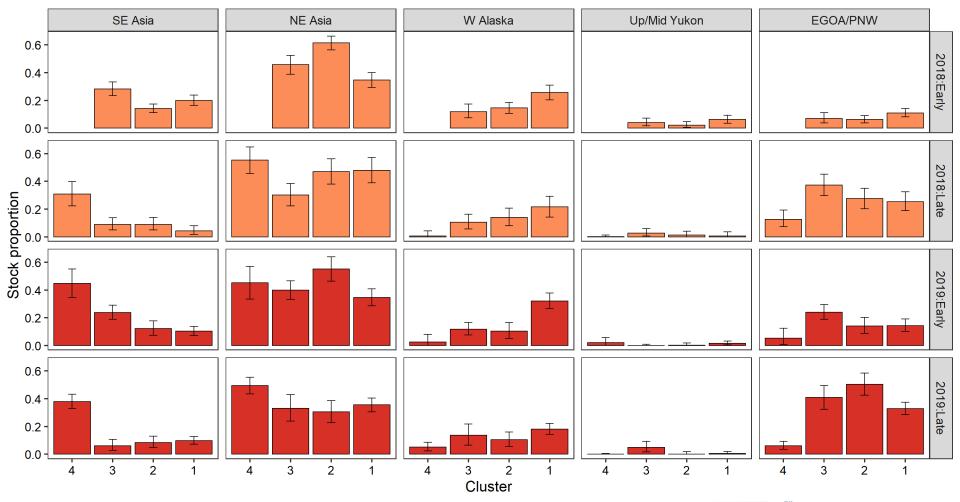
2019

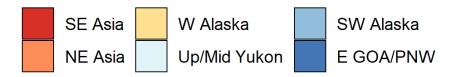


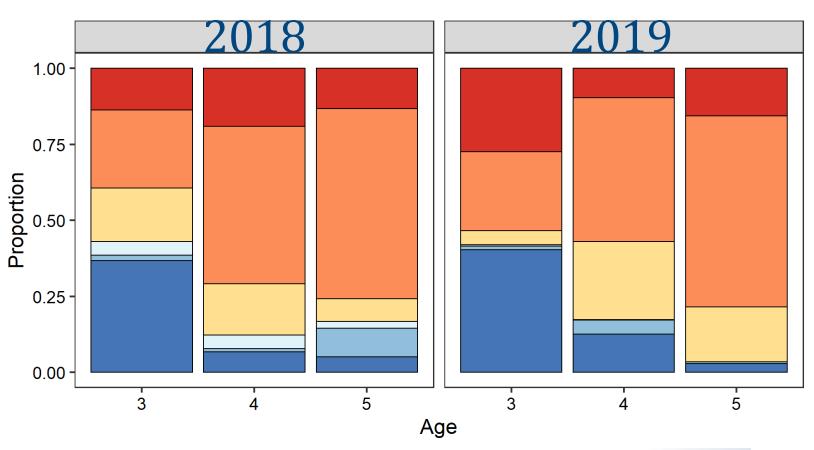




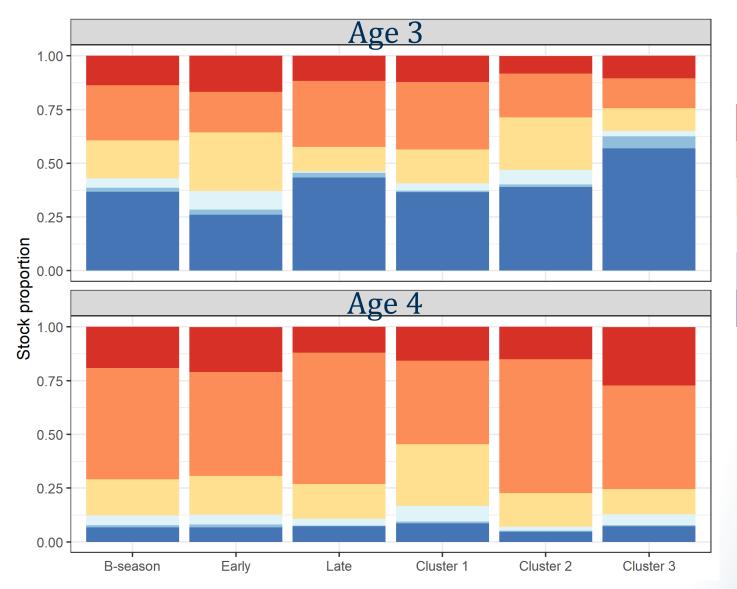










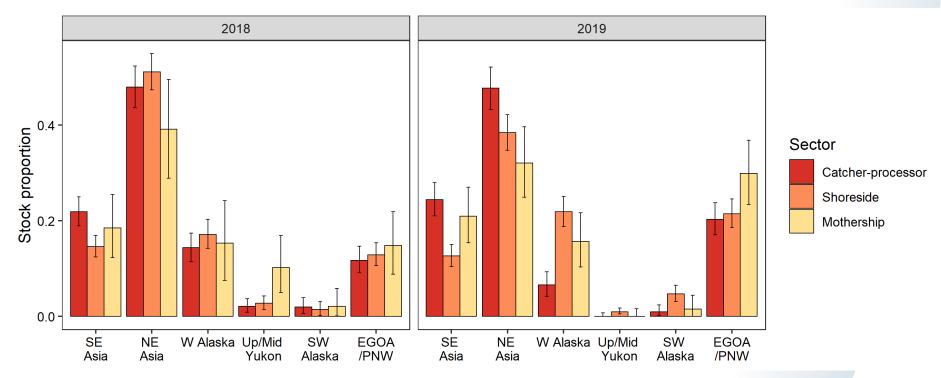






2018

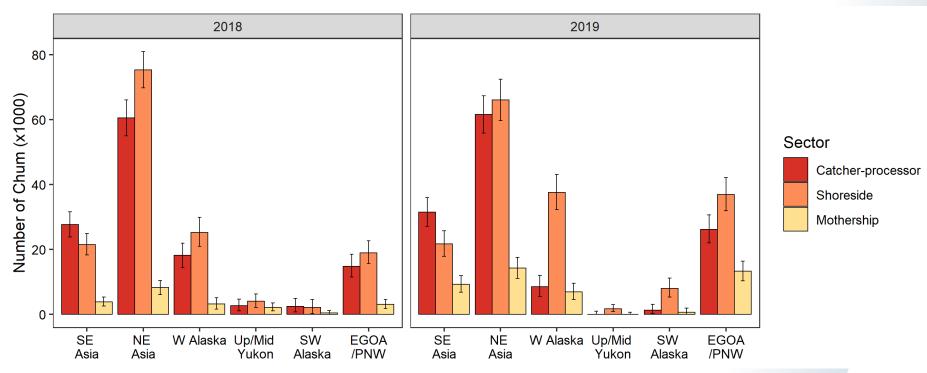
2019





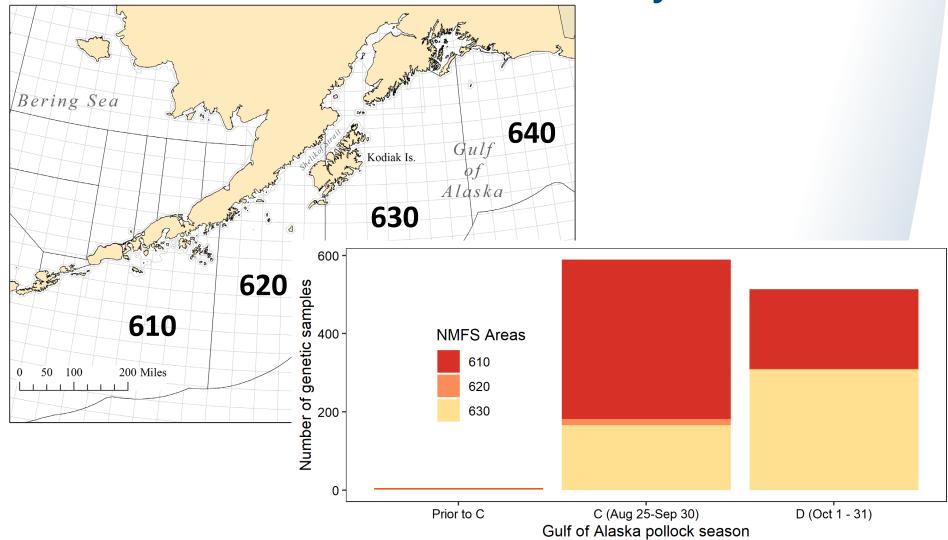


2019



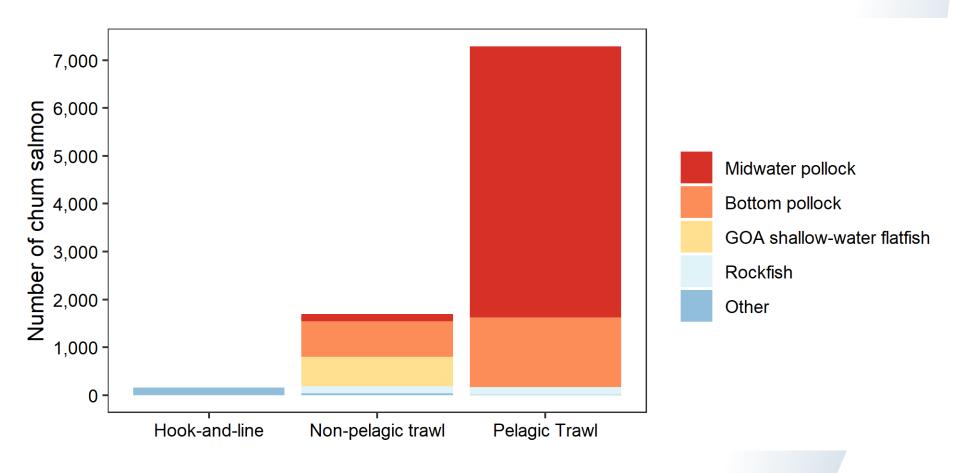


2018 GOA chum salmon bycatch



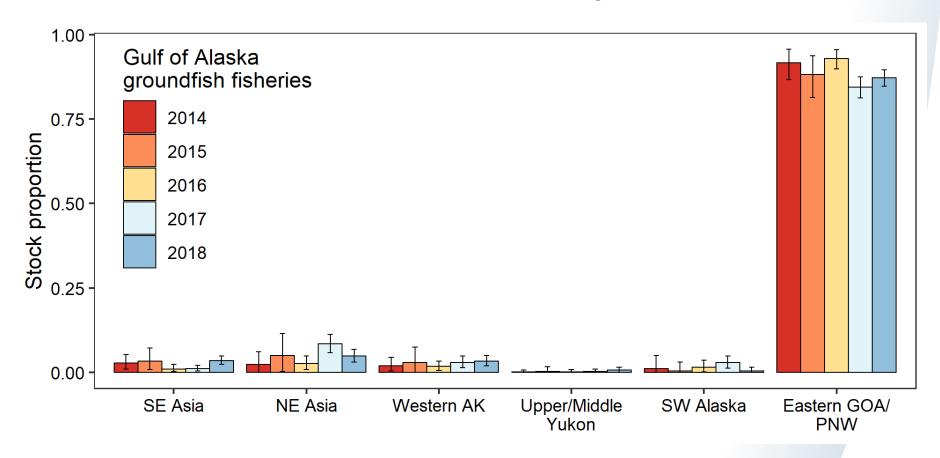


2018 GOA chum salmon bycatch





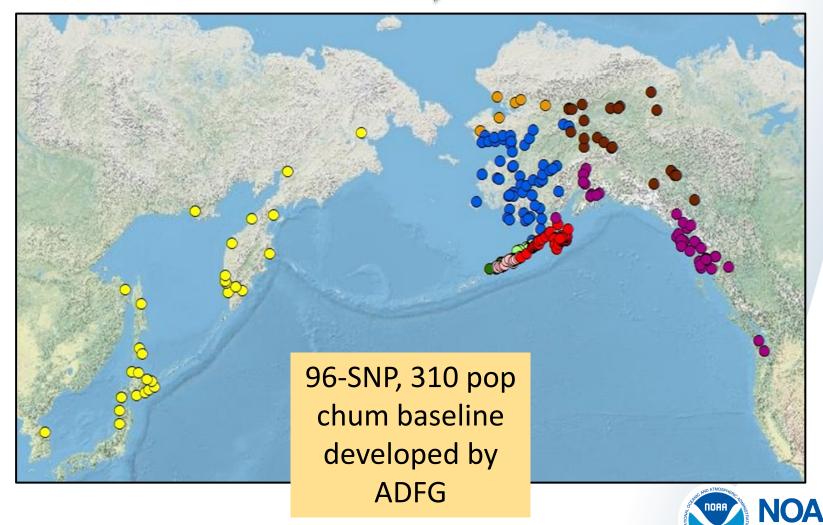
GOA chum salmon bycatch





2020 Bycatch

11 Microsatellites ———— 91 SNPs (GTseq)



Next steps

- 10 years of systematic sampling, genetic and age analyses behind us.
- New technologies, staff, PCCRC post-doc well-positioned to analyze trends.
- Over the next year, concerted effort to integrate:
 - ✓ spatial and temporal distributions of bycatch
 - ✓ genetics and age
 - ✓ fishing fleets/sectors
 - environmental information
- Improve stock-specific bycatch avoidance.





