

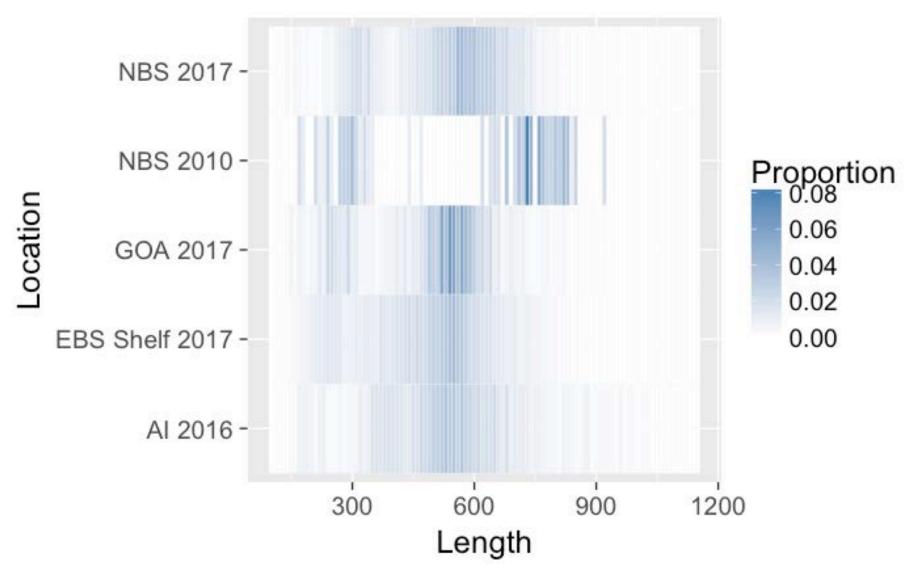


Proposed genetic analysis of Northern Bering Sea Pacific cod

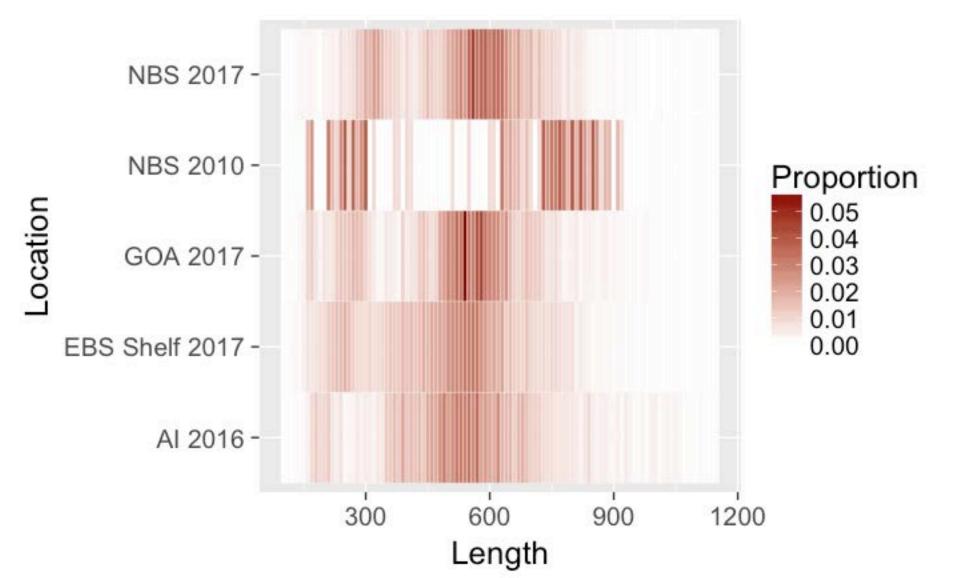
Ingrid Spies, Mike Canino, Dan Drinan, Lorenz Hauser



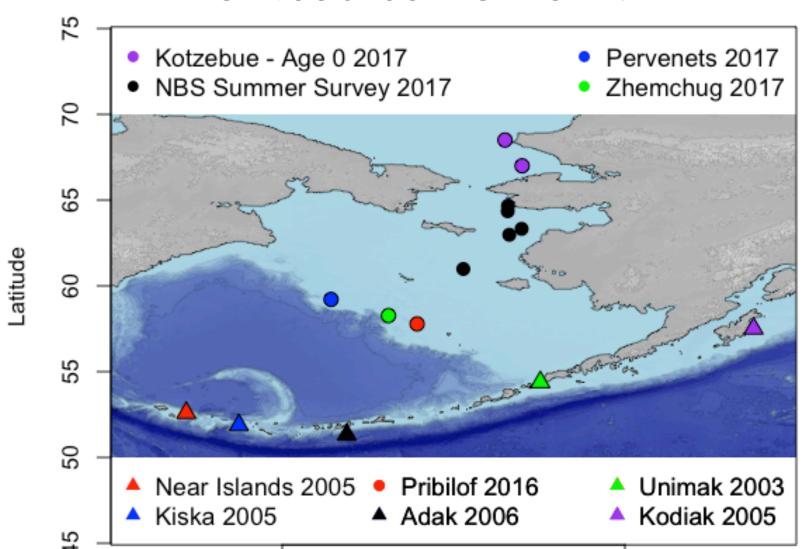
Length frequencies for Pacific cod in Alaska (males)



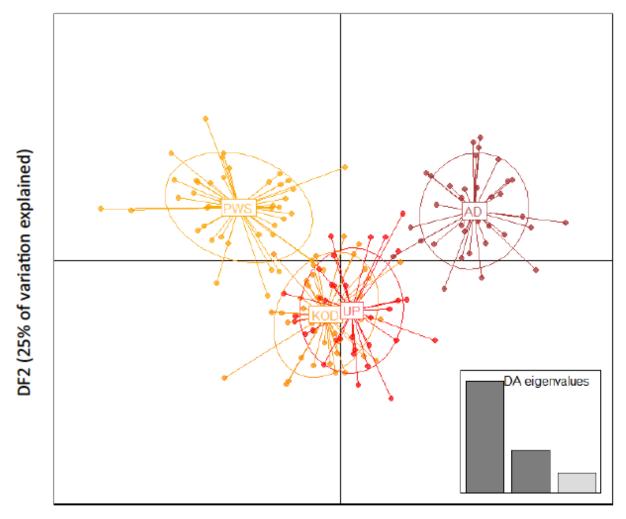
Length frequencies for Pacific cod in Alaska (females)



Pacific cod samples exist that can be used to examine: Where did all the NBS P.cod come from?



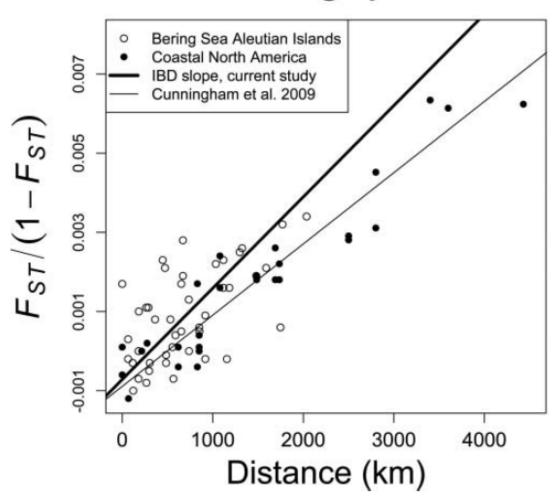
Loci have been found which can discriminate distinct spawning populations of Pacific cod in Alaska.



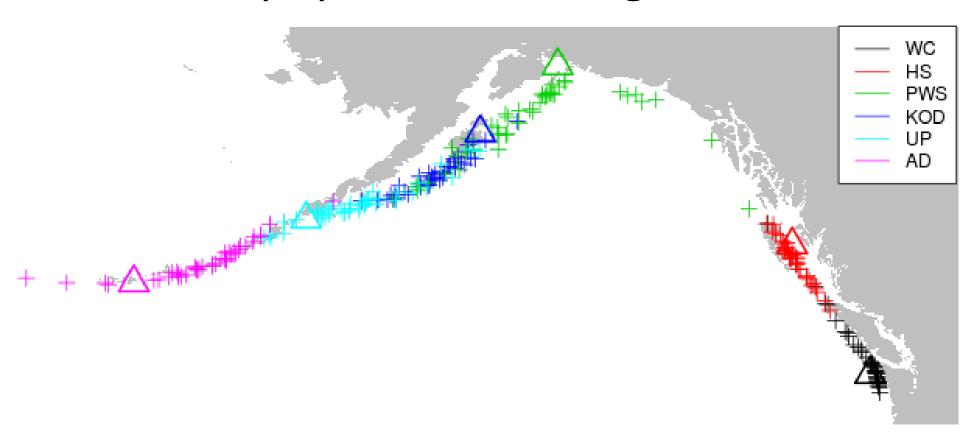
DF1 (64% of variation explained)

Isolation by distance pattern has been observed in Pacific cod

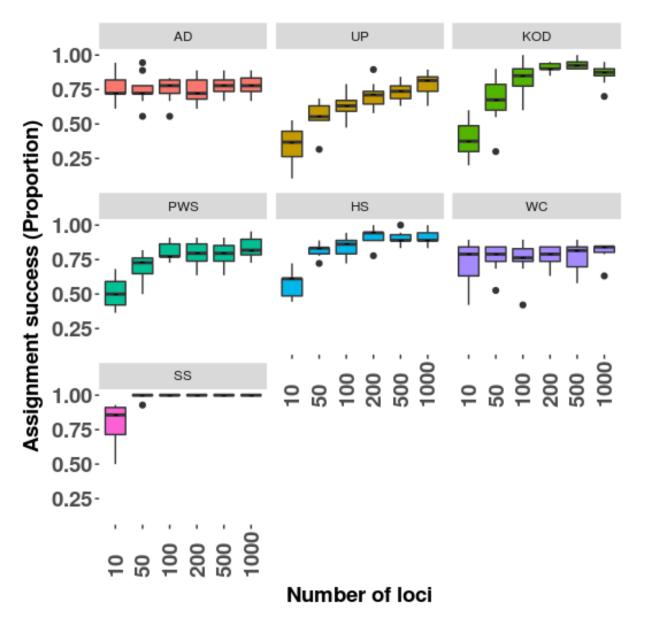
Genetic vs. Geographic distance



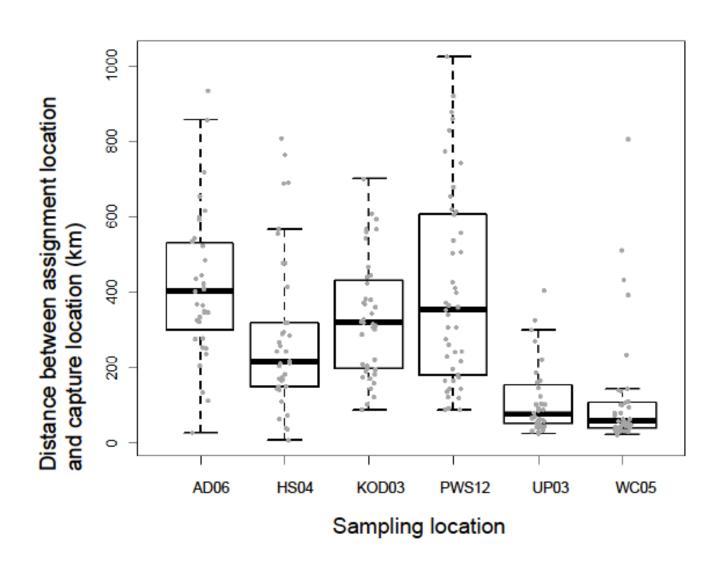
Genetic techniques have been developed to assign P. cod to their population of origin.



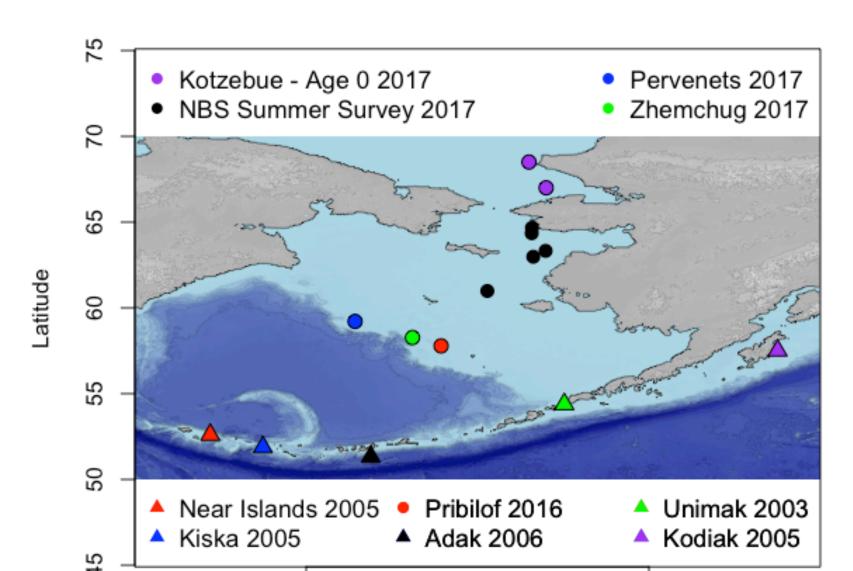
Assignment success depends on number of loci



Misassignment distance ~400km or less



Distance from Unimak Pass to Northern Bering Sea is ~1,000 km



Research questions relevant to management

- If NBS cod assign to stocks elsewhere, would they move back to those spawning areas in the winter?
- Have the cod in the Northern Bering Sea originated there?