

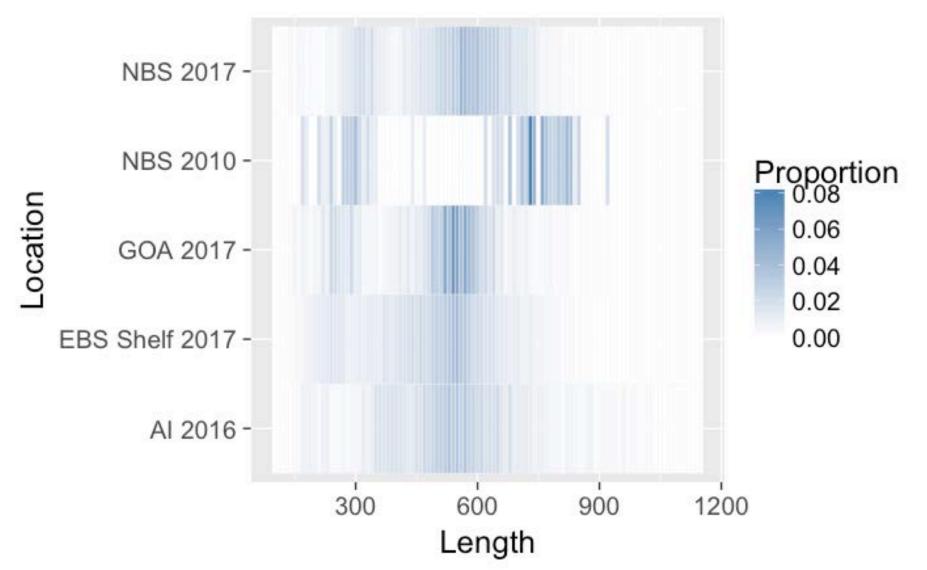


# Proposed genetic analysis of Northern Bering Sea Pacific cod

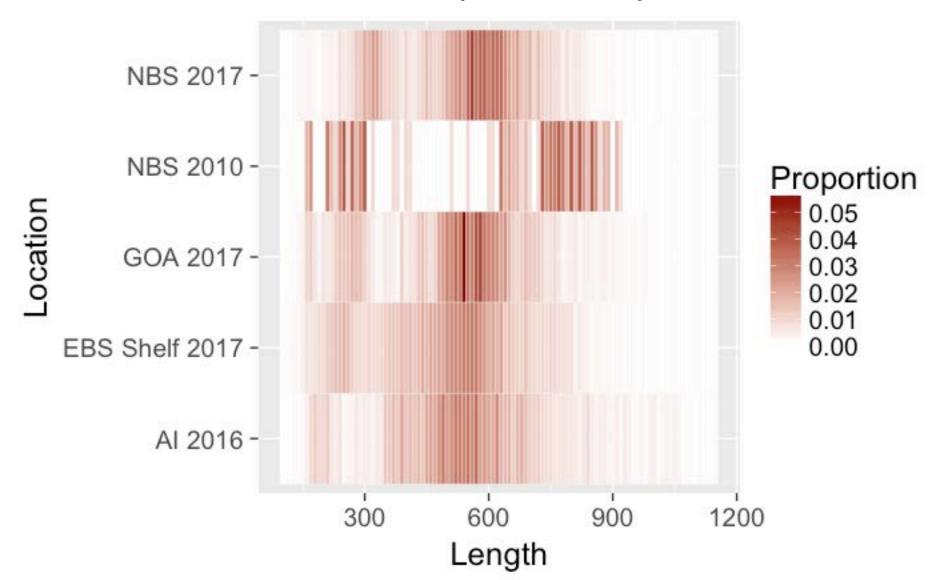
Ingrid Spies, Mike Canino, Dan Drinan, Lorenz Hauser



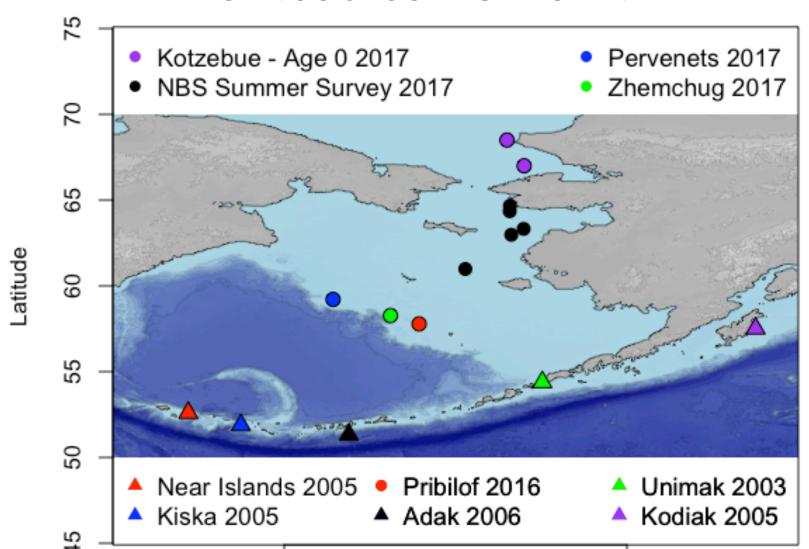
## Length frequencies for Pacific cod in Alaska (males)



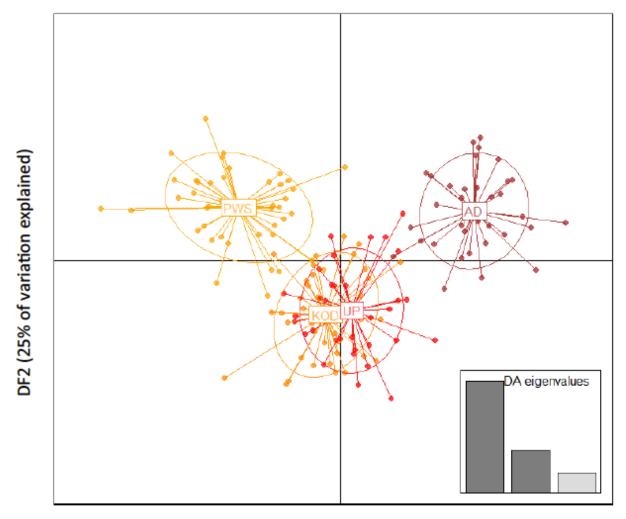
### Length frequencies for Pacific cod in Alaska (females)



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



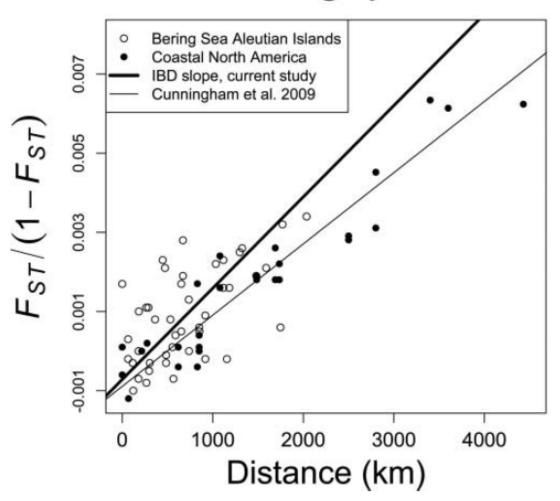
# Loci have been found which can discriminate distinct spawning populations of Pacific cod in Alaska.



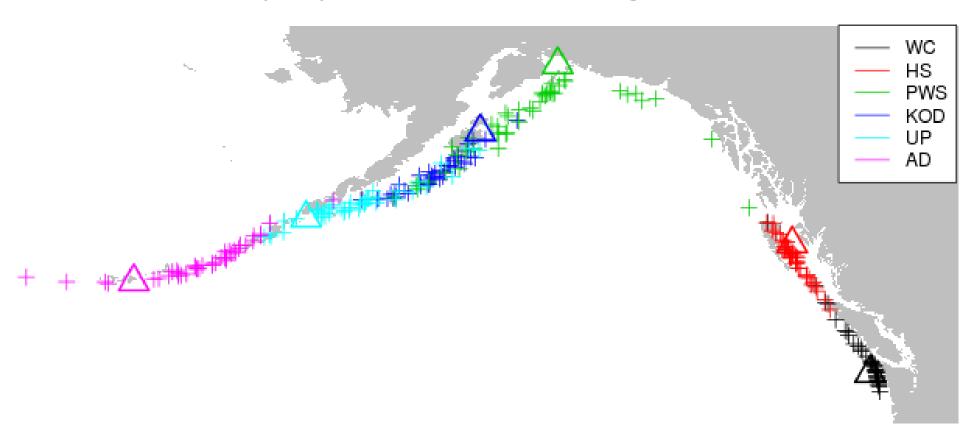
DF1 (64% of variation explained)

### Isolation by distance pattern has been observed in Pacific cod

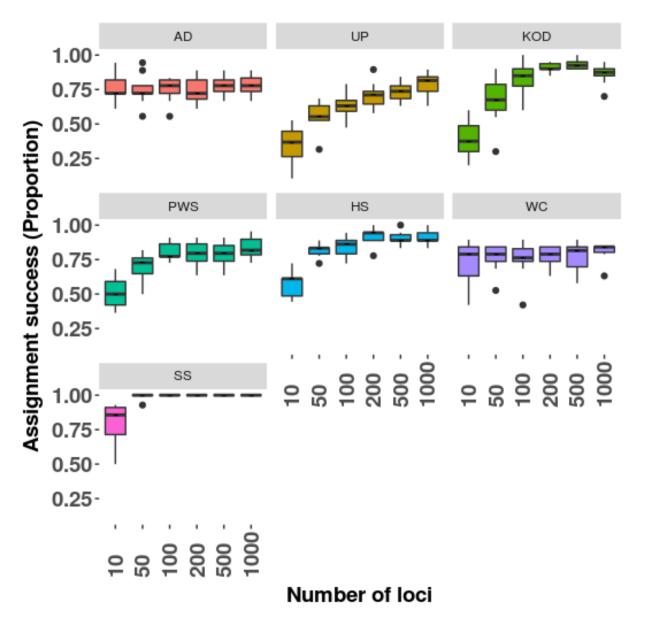
#### Genetic vs. Geographic distance



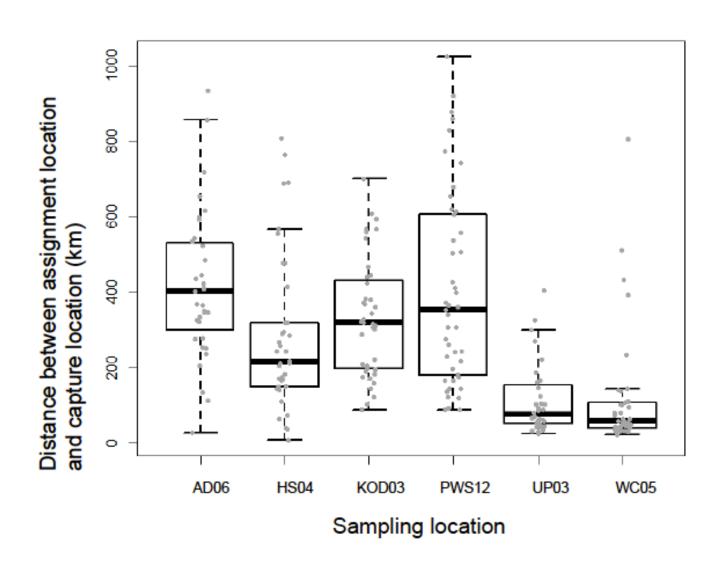
# Genetic techniques have been developed to assign P. cod to their population of origin.



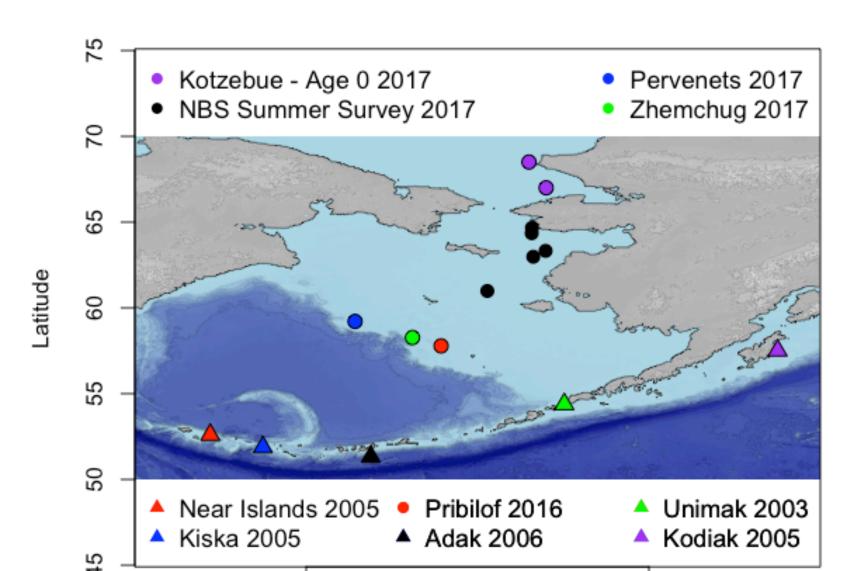
#### Assignment success depends on number of loci



#### 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?