### Department Framework for Interpretation of Results



C. Habicht and W. D. Templin

Alaska Department of Fish and Game Gene Conservation Lab Alaska Hatchery Research Program Informational Meeting March 7, 2019 AHRP identified three questions that would help inform policy and that were attainable

- 1) What is the genetic structure of pink and chum in PWS and SEAK?
- 2) What is the extent and annual variability of straying?
- 3) What is the impact on fitness (productivity) of natural pink and chum stocks?

Some questions that I've been asked that are not addressed by AHRP

- What are the competition and predation effects of hatchery fish?
  - Within and across species
  - Within marine and freshwater habitats
- Do hatchery fish reduce genetic resilience of wild populations?
- If changes in productivity are observed, what mechanisms could be driving these differences?
- How will findings affect policy?
- How do these hatchery fish in wild systems affect assessment of escapement?

### Department is assessing risk

- What we have now:
  - Wild system productivity
  - Hatchery proportions
- What we are working on now:
  - Contemporary population structure 90% PWS and SEAK
  - Historical population structure 50% PWS
  - RRS estimates 7% PWS, 0% SEAK
- Once all AHRG RRS results are complete:
  - RRS interpretation
  - Implications for assessment of escapement
- In the meantime, literature review
  - Genetic resilience of wild populations
  - Competition and predation effects of hatchery fish
    - Within and across species
    - Within marine and freshwater habitats
- Analyses and interpretation will inform policy maker decisions

### Conceptual model for assessing risk



### Conceptual model for assessing risk



### Conceptual model for assessing risk

Many generations (e.g. genetic)



#### Assessing mechanisms driving Relative Reproductive Success



C. Habicht and W. D. Templin

Alaska Department of Fish and Game Gene Conservation Lab Alaska Hatchery Research Program Informational Meeting March 7, 2019 RRS estimates: 7% complete RRS interpretation: 0% complete

- Inappropriate to interpret beyond:
  - 1 stream (Hogan Bay)
  - 1 generation for even- and odd-years
- Does not represent variation:
  - Across years, within stream
  - Across steams
  - Across generations (grandparents)
  - Across species (chum salmon)



### Example of RRS across years within species and location: Steelhead, Hood River





From Christie et al. 2014; original data Araki et al. 2007

# Examples of RRS across years within species and locations



From Christie et al. 2014; original data various sources

### RRS estimates: 7% complete RRS interpretation: 0% complete

- Inappropriate to interpret beyond:
  - 1 stream (Hogan Bay)
  - 1 generation for even- and odd-years
- Does not represent variation:
  - Across species (chum salmon)
  - Within stream, across years
  - Across steams
  - Across generations (grandparents)
- We do not know what is driving RRS
  - Once we have results, we can investigate mechanisms

Many mechanisms may drive measured RRS: Here are a few Many generations (e.g. genetic) One generation (e.g. non-genetic)

Relaxation of natural selection

### Relaxation of selection: a genetic example

- Hatcheries increase survival that's the whole point
- Most mortality in the wild is due to unsurvivable events, e.g.:
  - Too much rain scouring
  - Too little rain dewatering
  - Too cold freezing
  - Disturbance
- Some mortality in the wild is caused by genetic issues:
  - Most of these would die in a hatchery anyway
  - Some might survive in a hatchery, e.g.:
    - Lack of disease resistance
    - Inability to avoid predators
    - Tolerance of temperature or oxygen fluctuations
- The conditions in the hatchery do not select out the same fish as the conditions in the wild



Spawning ground familiarity: a non-genetic example

- Homing fish have the potential to find the location where they were incubated
- These incubation locations were suitable (otherwise the fish would not have survived)
- Staying fish (regardless of origin), need to identify a suitable location
- Straying fish that find suitable locations, produce progeny that, if they home, will have the homing fish advantage
- Straying fish that do not find a suitable location, will produce fewer (if any) progeny.
- Therefore, most of this effect is wiped out the next generation

Many mechanisms may drive measured RRS: Here are a few One generation Many generations (e.g. non-genetic) (e.g. genetic) Spawning ground familiarity Relaxation of natural selection Domestication selection **Epigenetics** Genetic drift Run timing-associated variables **Broodstock incompatibility** 

Sexual selection

Fishery prosecution

- Spawning ground competition
- Straying fish delays

Data available to investigate mechanisms driving RRS

- Genetic mechanisms
  - Modeling
  - Grandparent RRS
  - Historical and contemporary genetic structure (PWS)
- Non-genetic mechanisms
  - Timing of spawning
  - Location within stream
  - Fishery prosecution



## Questions?