

Genetic effects of Straying

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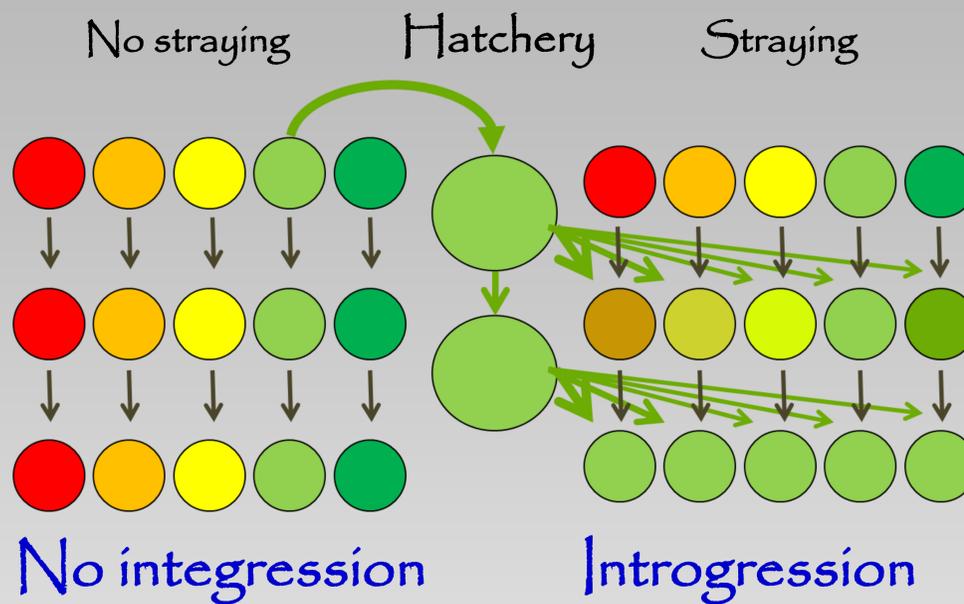


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Why worry about hatchery strays?

- Hatchery culture can shift the genetic makeup of fish¹
- Introgression reduces adaptive fitness of wild fish
 - Quantitative genetic model predicts decline in fitness²
 - Shift in run timing with persistent straying³
- Strays can influence wild populations ecologically
 - Straying can reduce effective size of wild population
 - Compete for mates.
 - Compete for spawning sites.
 - Hybrid offspring compete with wild offspring



Genetic effects of straying

- Colonize new habitats
- Hybridization may produce mal-adapted offspring
- Straying can lead to genetic swamping (Ryman-Laikre effect)⁴
- Introgression can lead to reduced fitness of wild population.
- Persistent straying reduces genetic diversity among wild populations.

What influences straying?

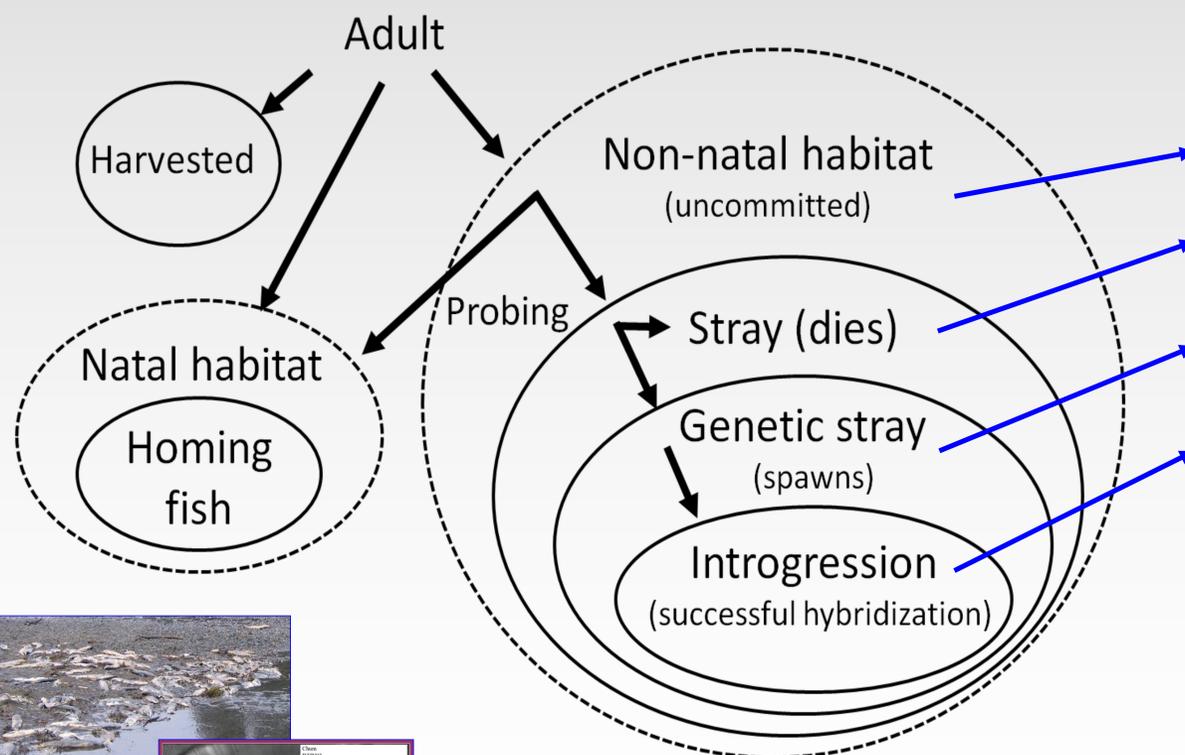
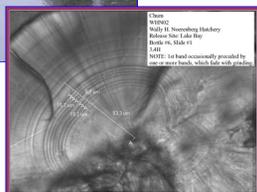
- Life history differences among species
- Size of hatchery release
- Method of releasing & imprinting
 - remote releases
 - freshwater-salt water transition
- Distance from release site

How to identify a stray?

- Physical marks—CWT, thermal
- Genetic marks
- In-season identification is difficult.



Photos: Rich Brenner



Not all 'stray' fish spawn

- Some fish probe into but will eventually home to their natal area
- Some fish die before spawning—naturally or by predation
- Some fish spawn with wild fish to produce hybrids.
- Introgression occurs only when hybrids survive, return and spawn

References:

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4. Araki H *et al.* 2008 Fitness of hatchery-reared salmonids in the wild. *Evol. Appl.* 1: 342-355.
5. Ford MJ 2002 Selection in captivity during supportive breeding may reduce fitness in the wild. *Cons. Biol.* 16: 815-825.
6. Ford MJ *et al.* 2006 Changes in run timing and natural smolt production in a naturally spawning coho salmon population after 60 years of intensive hatchery supplementation. *Can. J. Fish. Aquat. Sci.* 63: 2343-2355.
7. Ryman N & Laikre L 1991 Effects of supportive breeding on the genetically effective population size. *Cons. Biol.* 5: 325-329.