

# Migratory *O. mykiss* Colonizing a Natal Stream After Barrier Removal in the Methow River basin, WA

Dana Weigel, Univ. of Idaho, Dept. Fisheries and Wildlife,  
Patrick Connolly, USGS, Columbia River Research Lab and  
Madison Powell, Univ. of Idaho, Aquaculture Research Institute

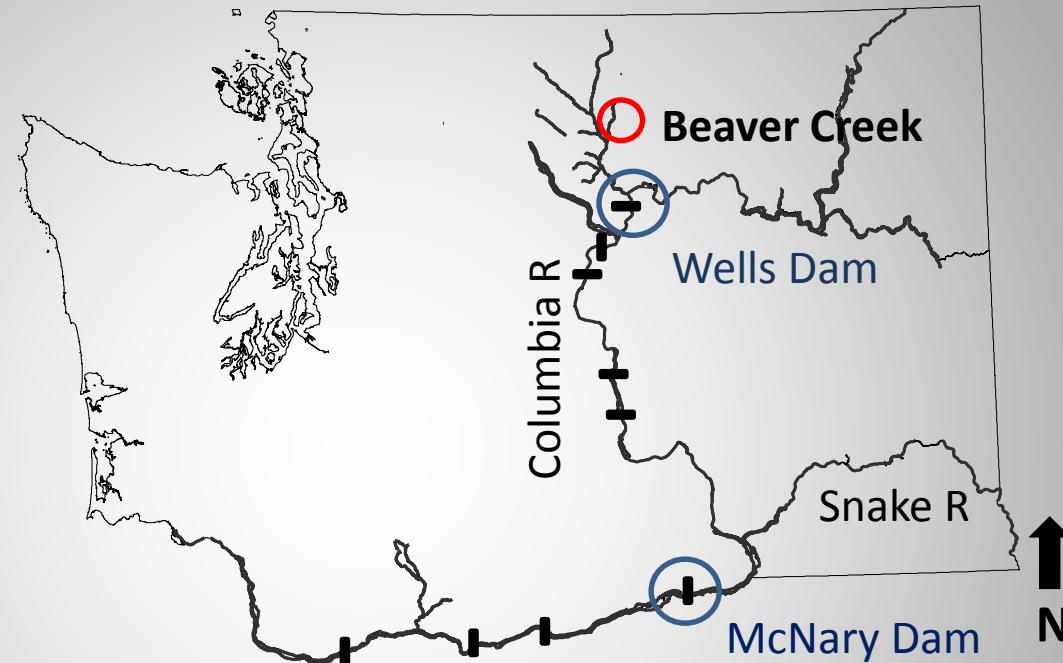


Willamette Science Review  
February 2017

# Study Area – tributary Methow River, upper Columbia Basin

Long mainstem migration (843 km)

9 mainstem dams

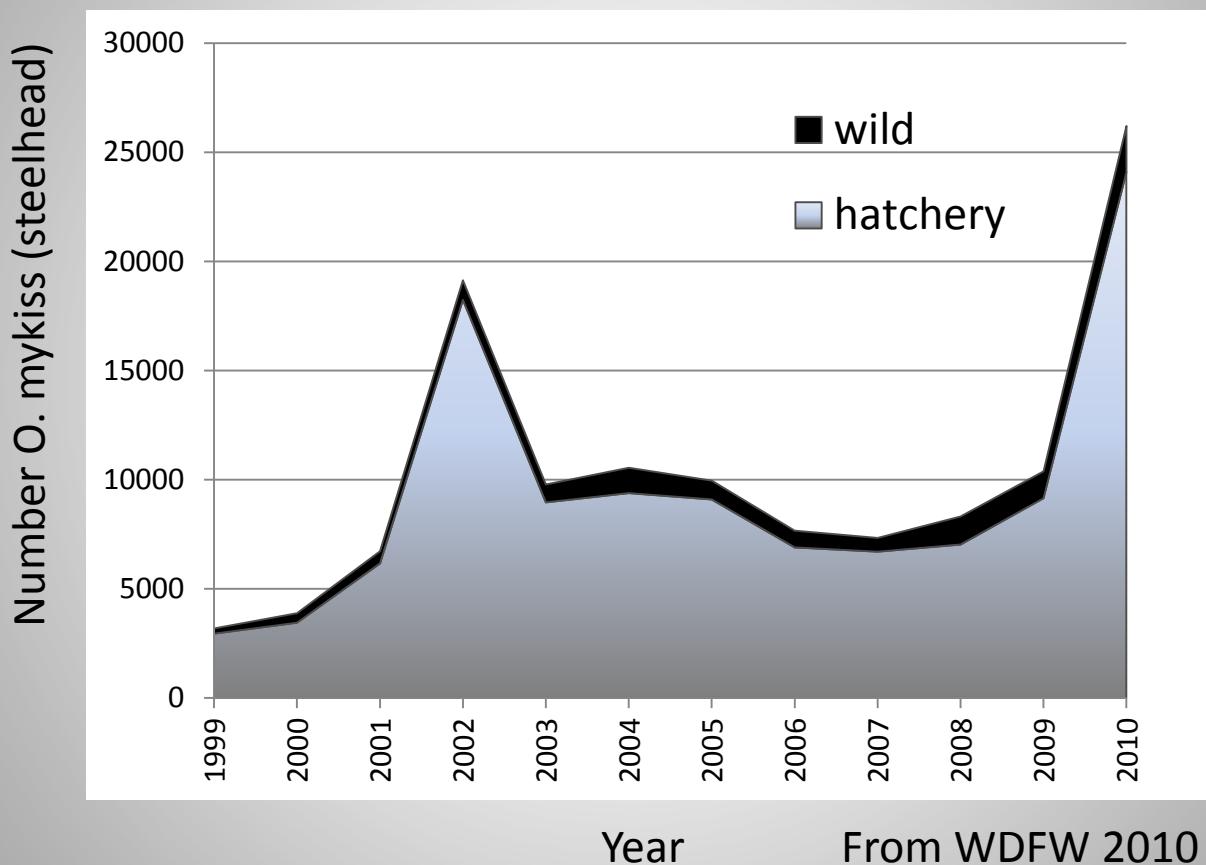


Upper Columbia Steelhead listed endangered 1997 -> threatened

Abundance	<5% extinction risk 100 yrs	
Productivity	3,000 spawners	2010 Status Review Not viable
Spatial Structure	Previously occupied habitats	
Diversity	Natural patterns genetic, phenotypic diversity	

# Study Area - Steelhead returns to Wells Dam 1999-2010

80-90% returning adults hatchery-origin



# Study Area - 7 Diversion dams redesigned in Beaver Creek 2002-2004



# Objectives

- Do migratory steelhead establish a population in the re-opened habitat?
- Which source -> establishes a population?
  - Individual reproductive success
  - mate selection, successful phenotypes to next generation
  - Founder effect? Or low diversity
- What spatial extent of colonization 1 generation after re-open habitat?
  - Migration strategy (tag tracking)
  - Genetic

# Three major processes shaping genetic structure ~

## Gene Flow ~

Migration  
Decreases genetic  
Differences thru  
mixing

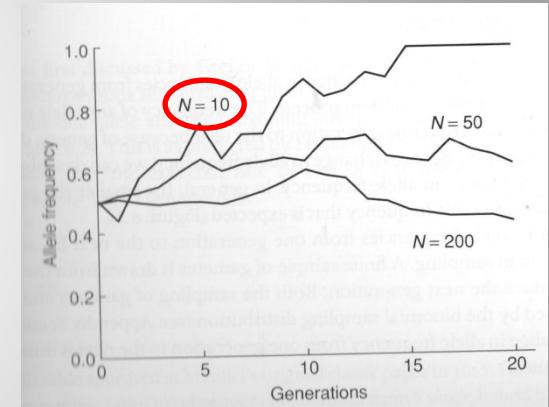
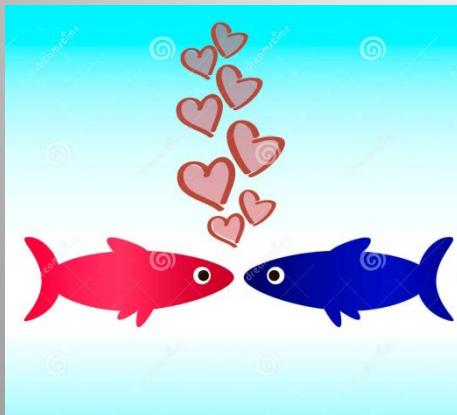


## Drift ~

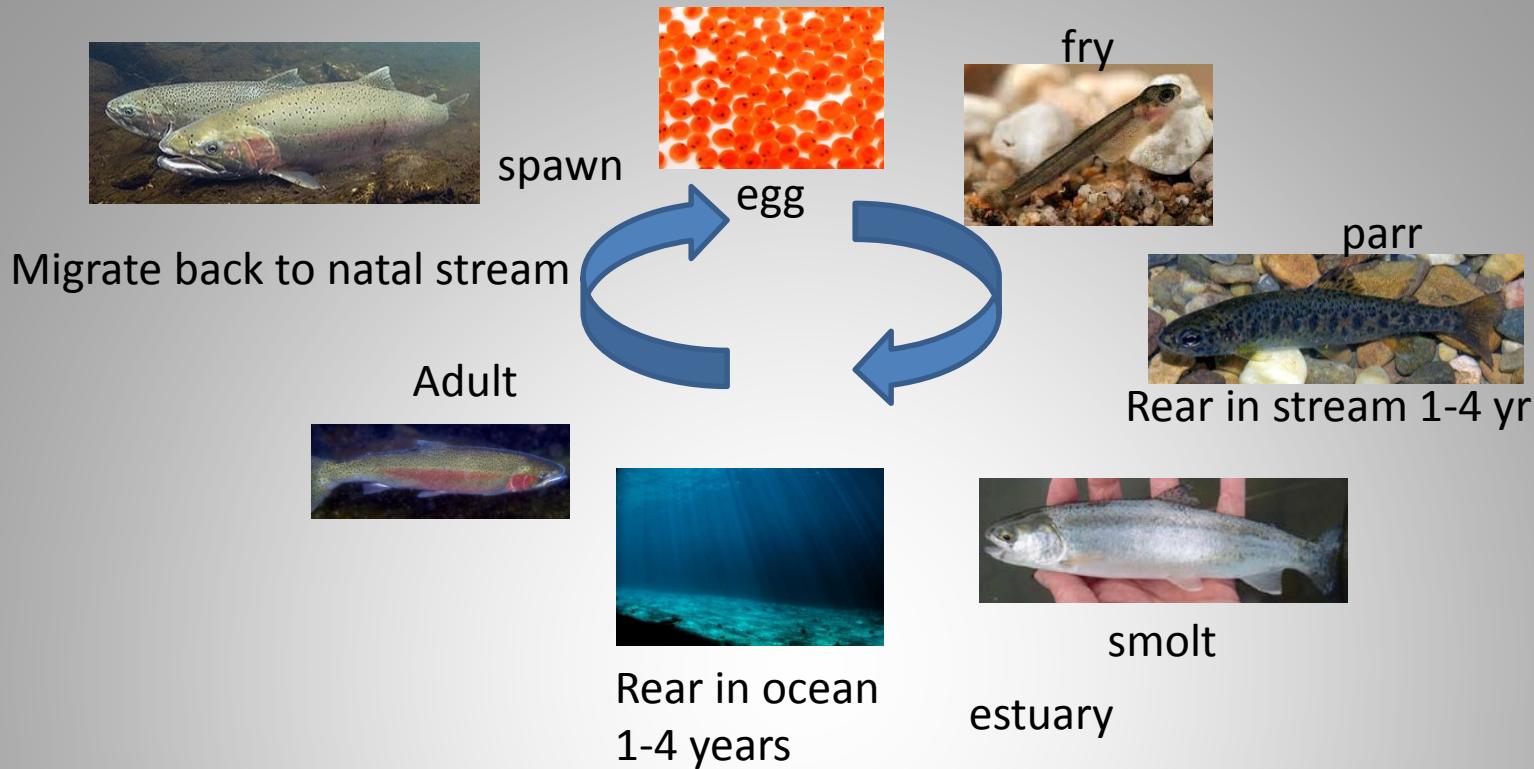
Increases genetic  
Differences  
Random fixation alleles  
Due to finite popn size

## System of Mating ~

selection for trait – genetic link



## Mutation Selection



## Life history diversity

### Migration

Anadromous  
Estuary  
Fluvial (River)  
Resident (Stream)  
Adfluvial (Lake)

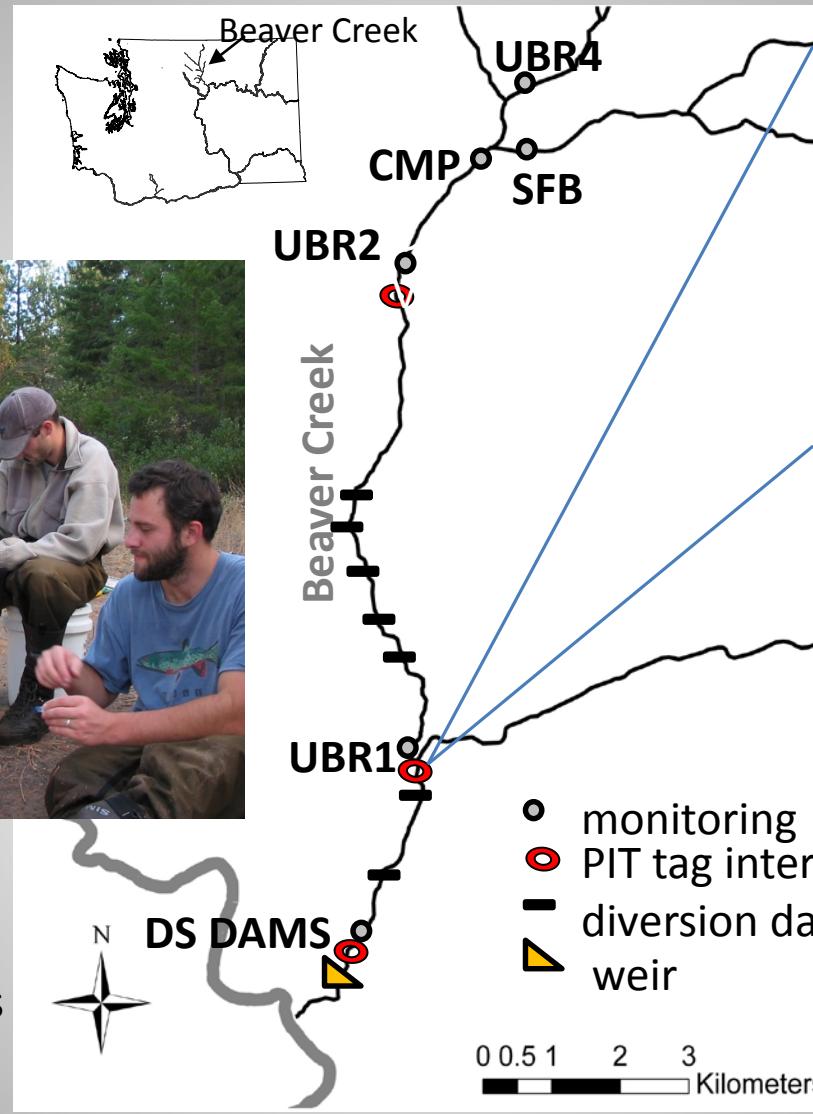
Steelhead

Rainbow Trout

Oviparous  
Overlapping generations

### Fresh:salt

1:1	2:1	3:1
1:2	2:2	3:2
1:3	2:3	3:3
1:4	2:4	3:4



Near mouth Methow  
Dams on Columbia R



# Methods – Capture

- Upstream/downstream picket weir
  - Feb to Dec 2005, 2006, 2007, 2008
  - Adults on spawning migration
  - Parr outmigrating

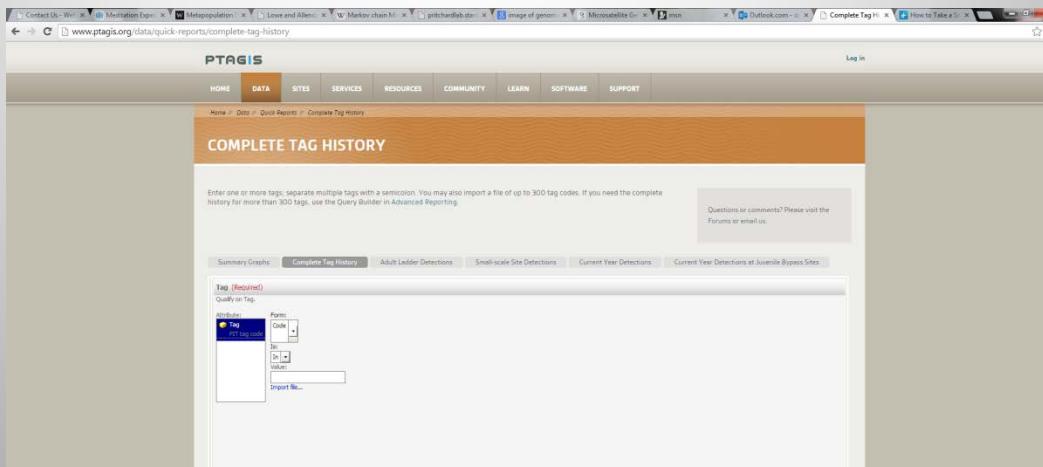


- Electrofishing at monitoring sites
  - Juvenile rearing in tributary



# Methods - Tagging

- Migration behavior was identified using PIT tag
- 16 digit alpha-numeric code
  - 3D9.1BF1FDC829
- Followed tags of juveniles reared in Beaver Creek to return as adults
  - PTAGIS



# Methods - Sample Selection

- Used Spawners from BY 2005 and 2006
- Queried tissue samples for analysis by length to match these brood years collected 2005-2008

Collection/BY age	2005	2006	2007	2008
2005	0			
2006	1	0		
2007	2	1	0	
2008	3	2	1	0

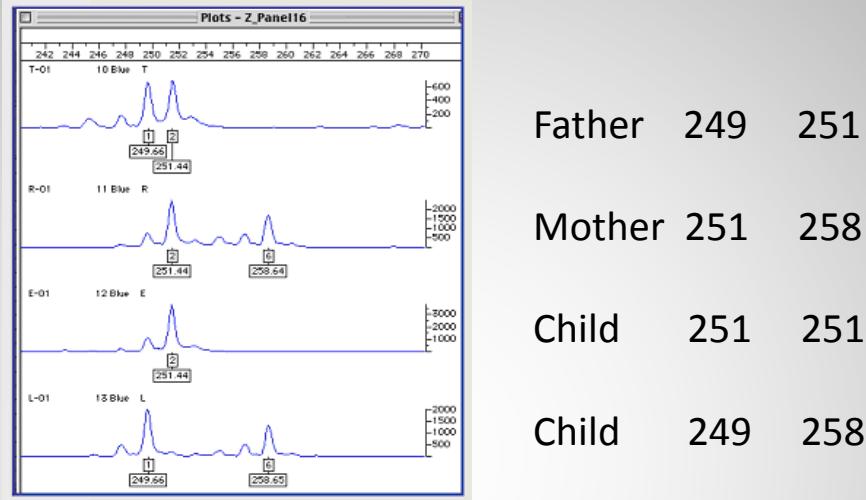
Collected Wells Hatchery tissue from WDFW for reference

# Methods - Tissue collection and Genotyping

- Fin clip preserved in 95% EtOH
- DNA extracted
- PCR amplification optimized for each locus
- **16 usat loci** – analysis conducted at UI Aquaculture Research Station, Hagerman ID
  - 13 usat loci standardized (Stevensen et al. 2009)
  - *One102* (Olsen et al 2000), *Omm1036* and *Omm1046* (Rexroad et al 2002)

# Parent – Offspring Matching

Exclusion tests with **1 mismatch** using Cervus > match 15 of 16 loci



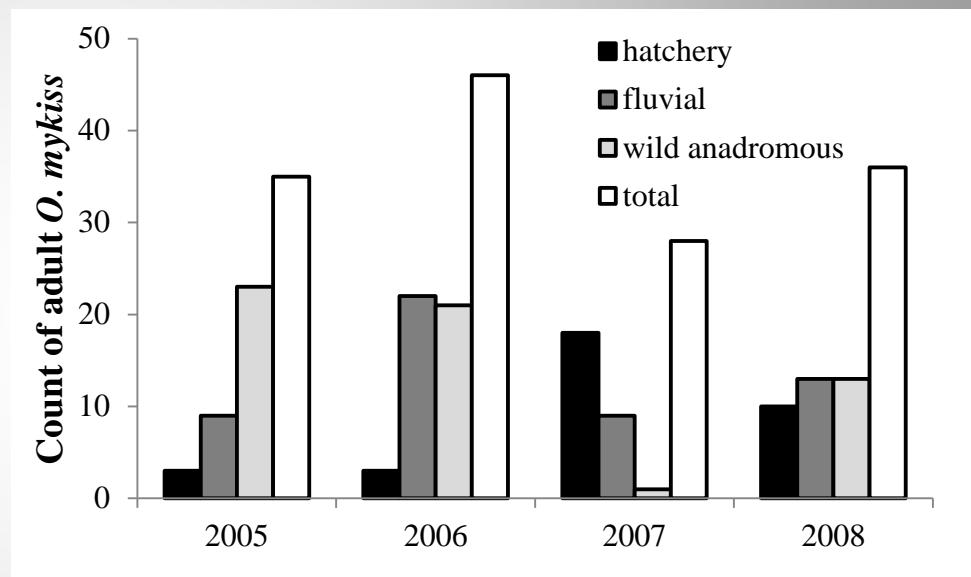
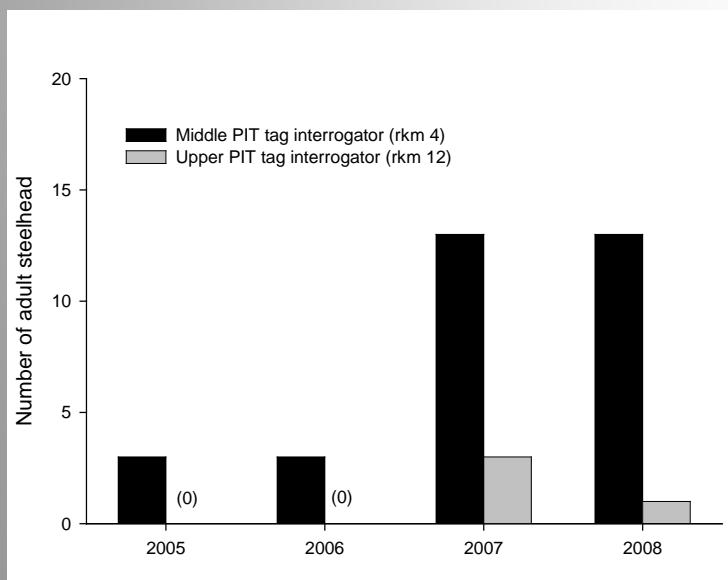
## Relate Successful Reproduction to Mate Selection

- Use Spearman Rank correlation to look at association between mating pairs

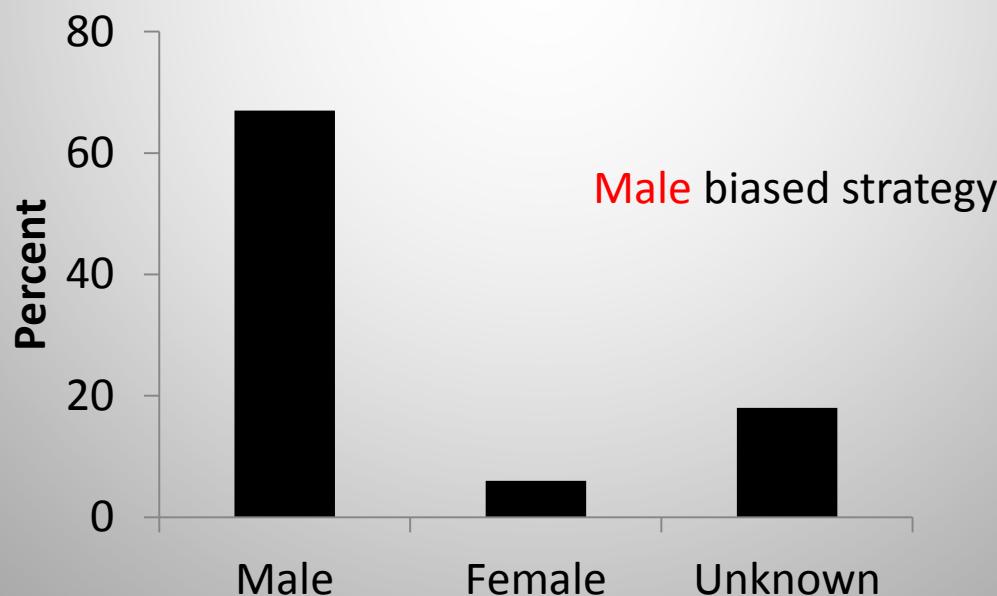
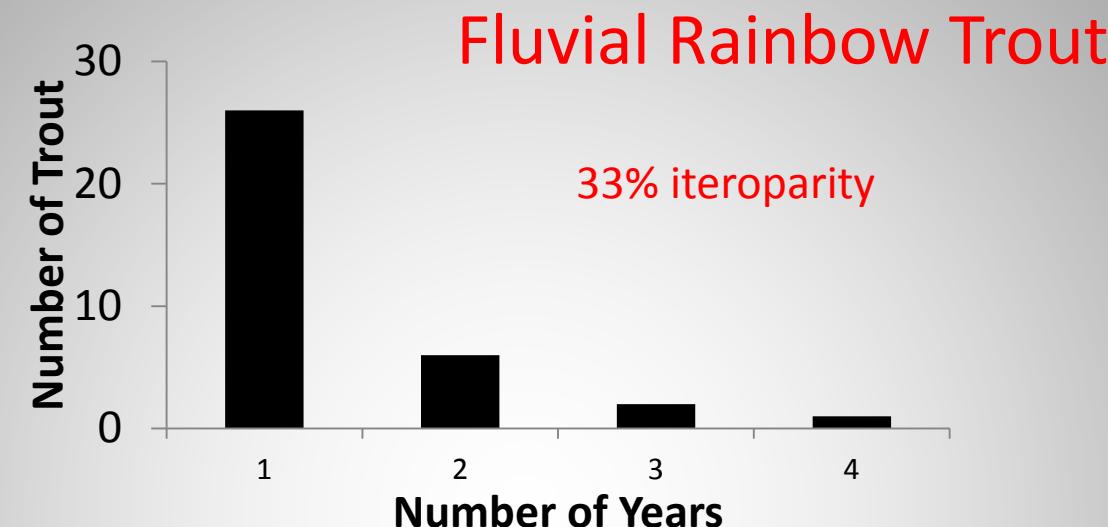
## Change in Popn Genetics over Generation

- $F_{ST}$  and Fisher exact tests

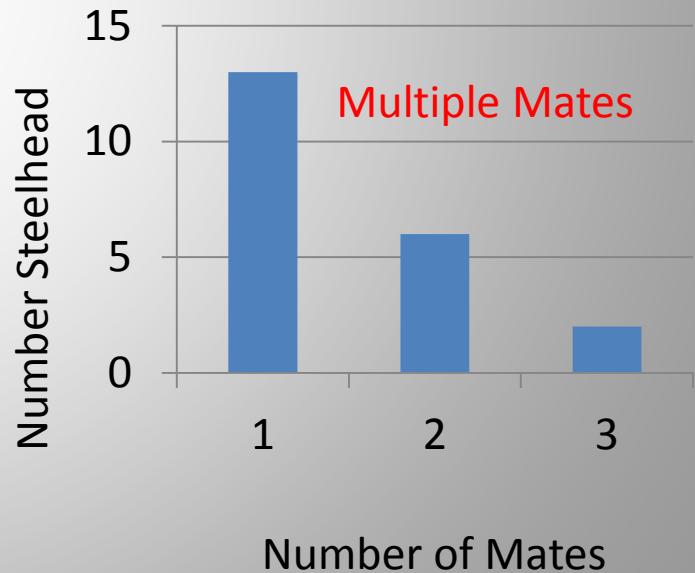
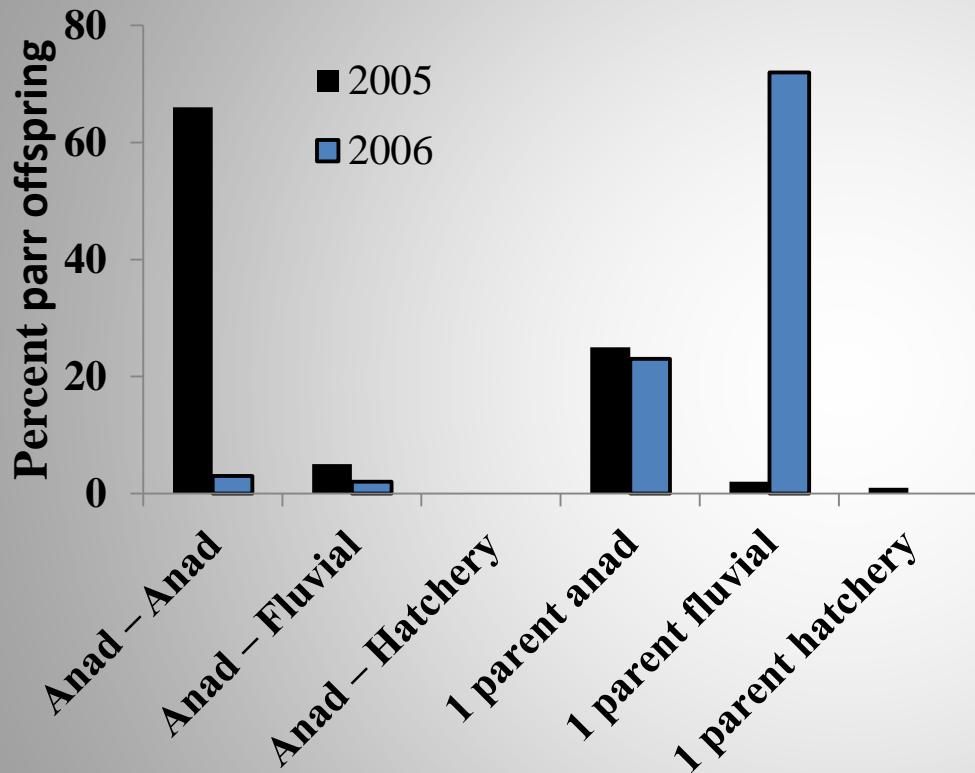
# Results – Do steelhead enter the re-opened Habitat? Source?



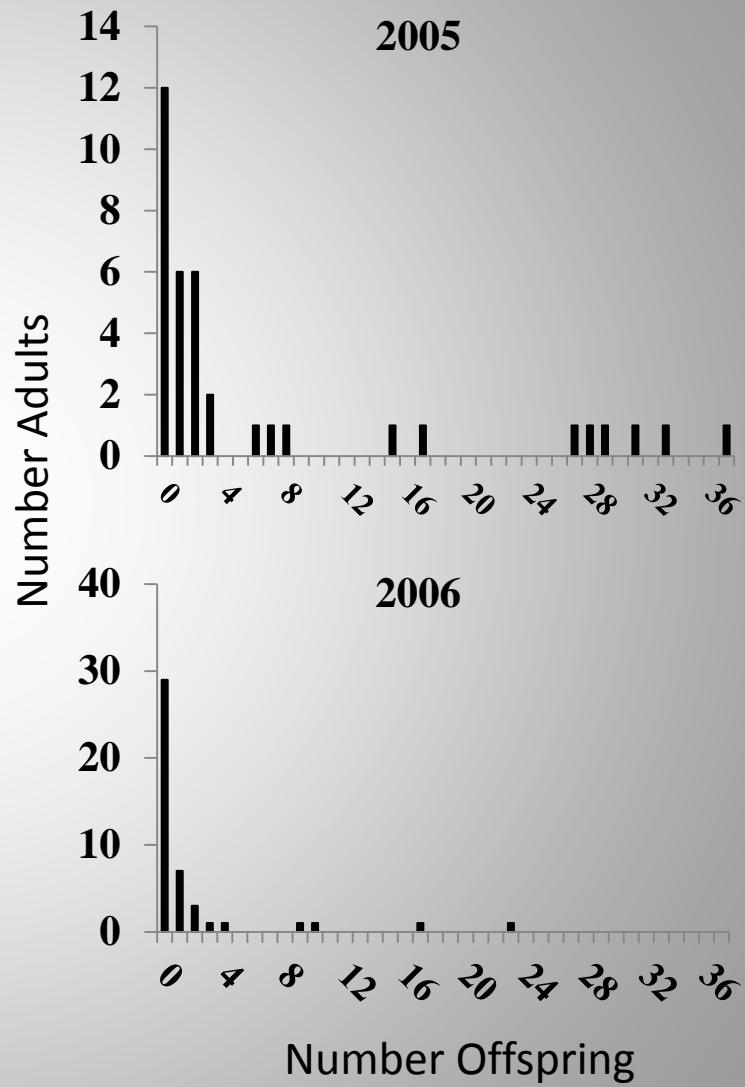
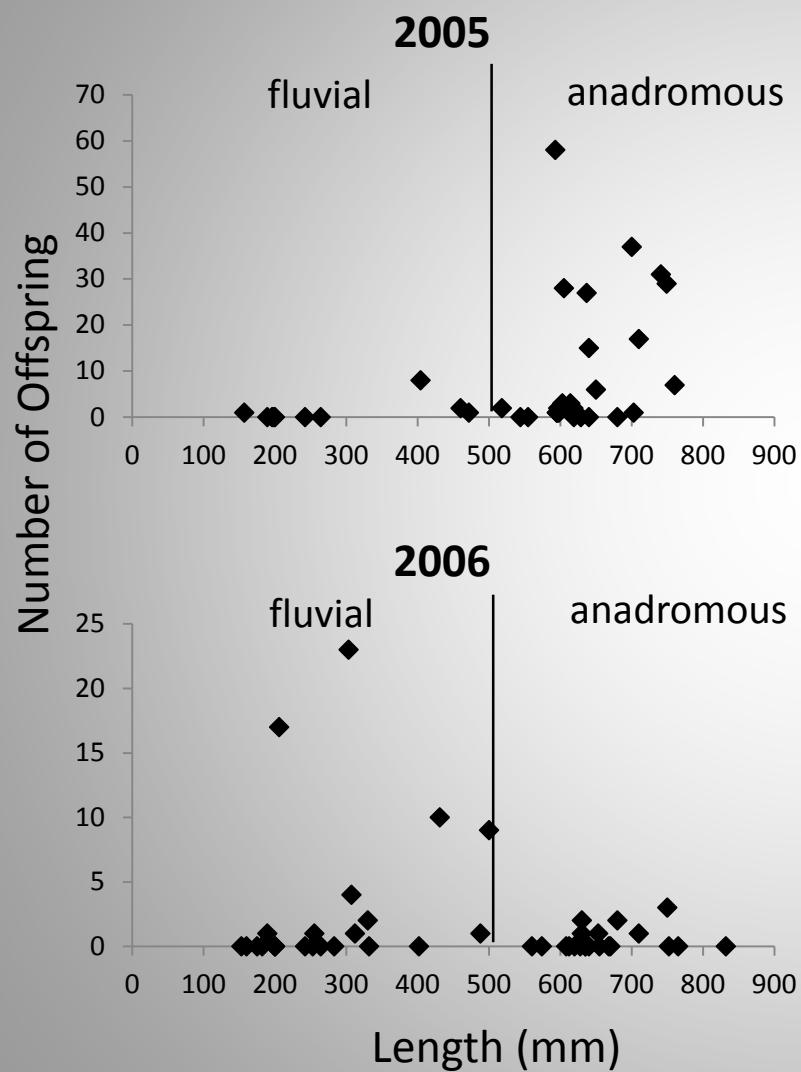
# Results – Do steelhead colonize the re-opened Habitat? Source?



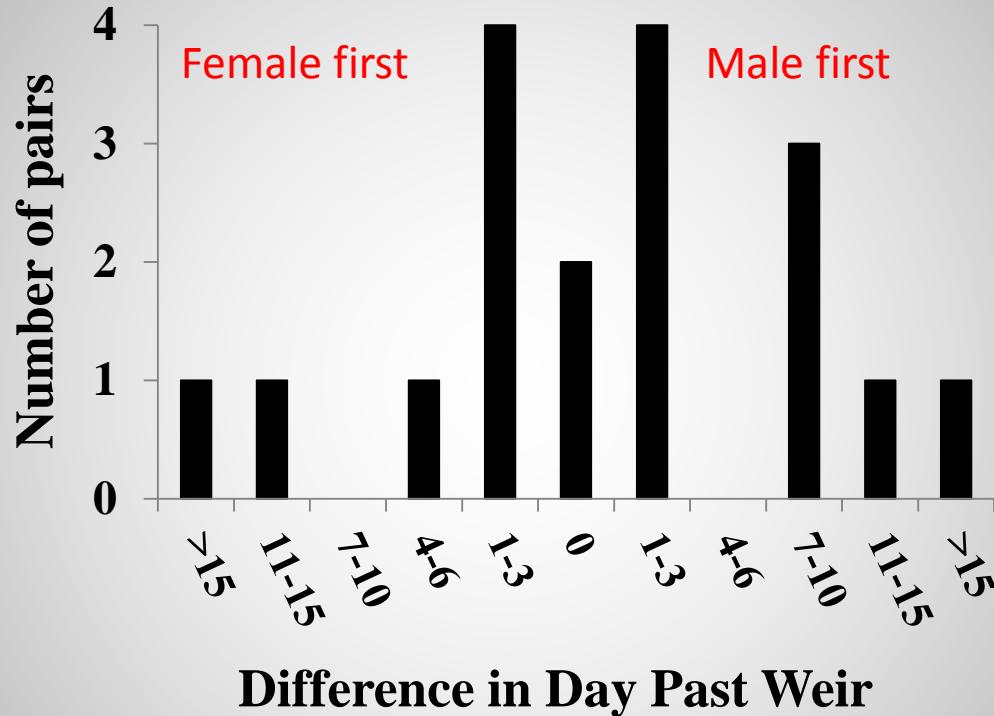
## Results – Successful mate selection by source?



# Results –Successful spawners by size/life history

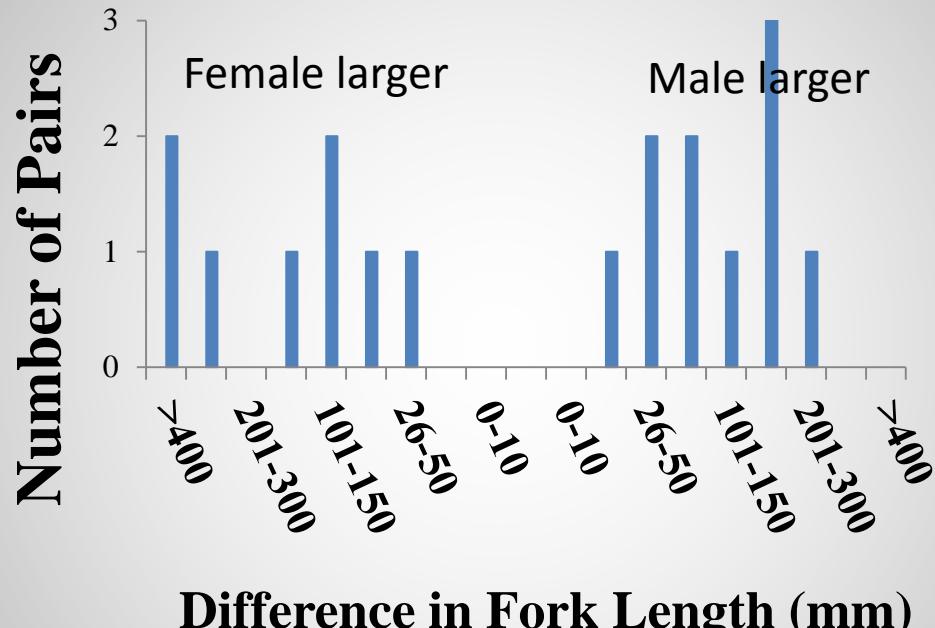


# Results - What phenotypes were associated with successful mating?



- Sig. paired by day past weir (early-early, later-later) ( $p<0.001$ ,  $\rho=0.84$ )

# Results – What phenotypes were associated with successful mating?

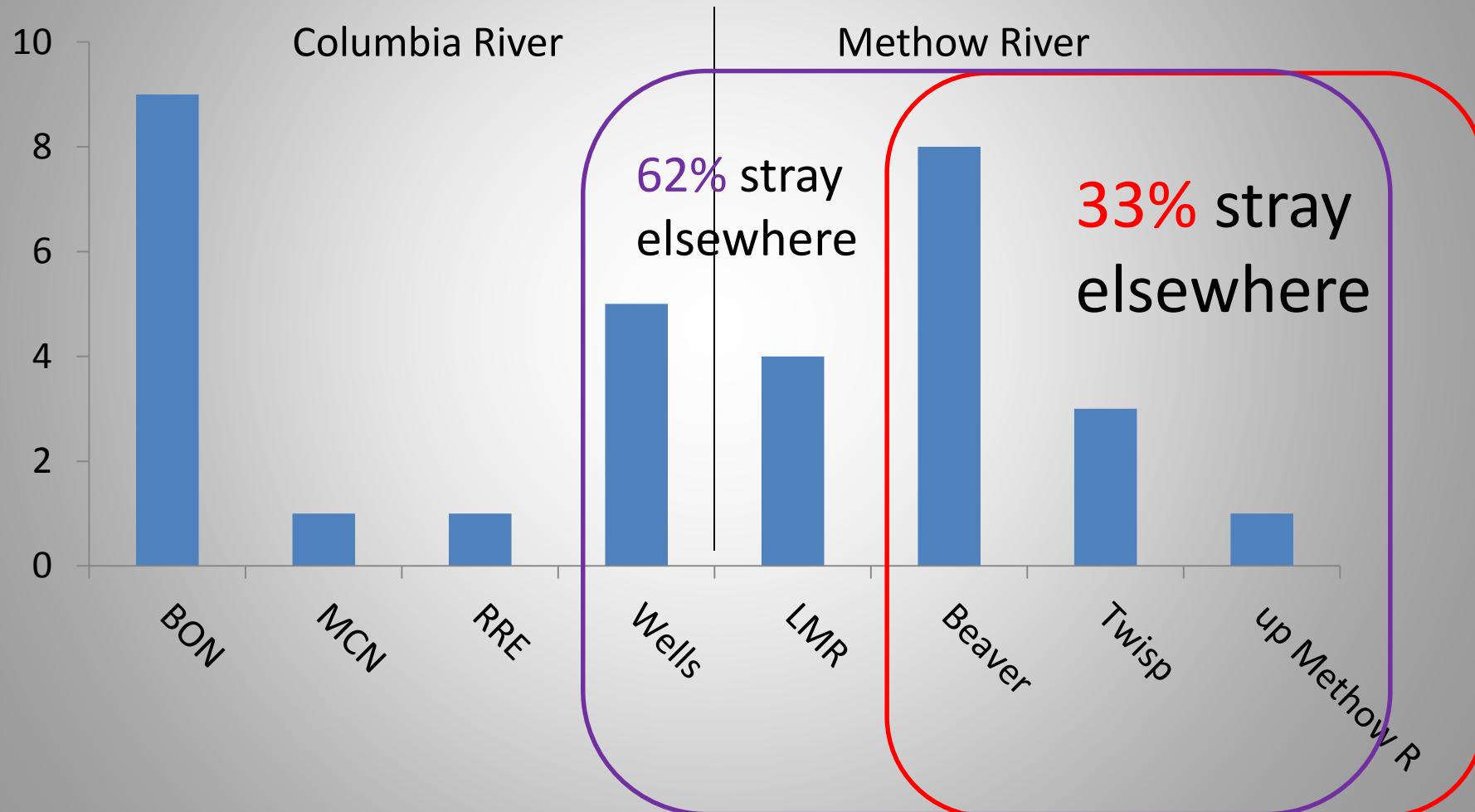


Sig. disassociation in size (larger-smaller) ( $p=0.005$ ,  $\rho=-0.72$ )

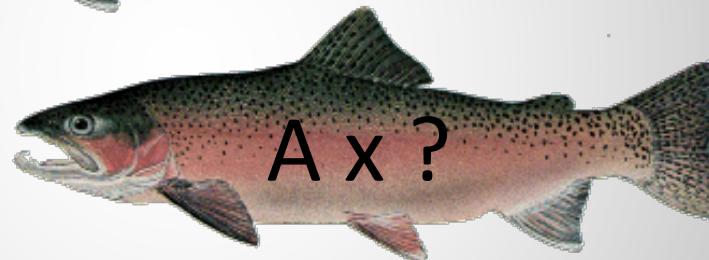
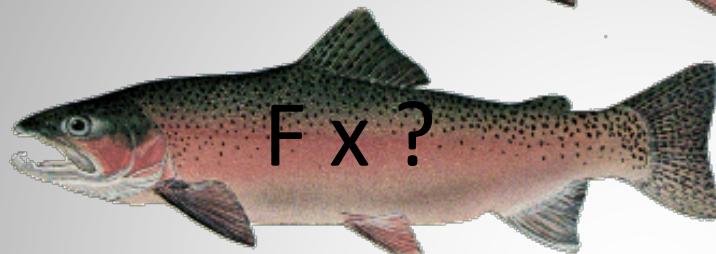
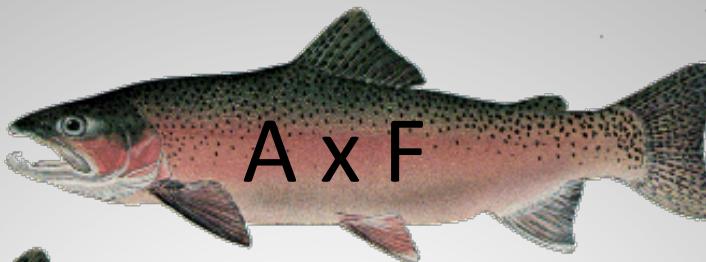
High genetic exchange among life history and generations

Random mating => no founder effect

# Results - Do steelhead colonize the reopened habitat?



# Results - Returning Offspring as Adults



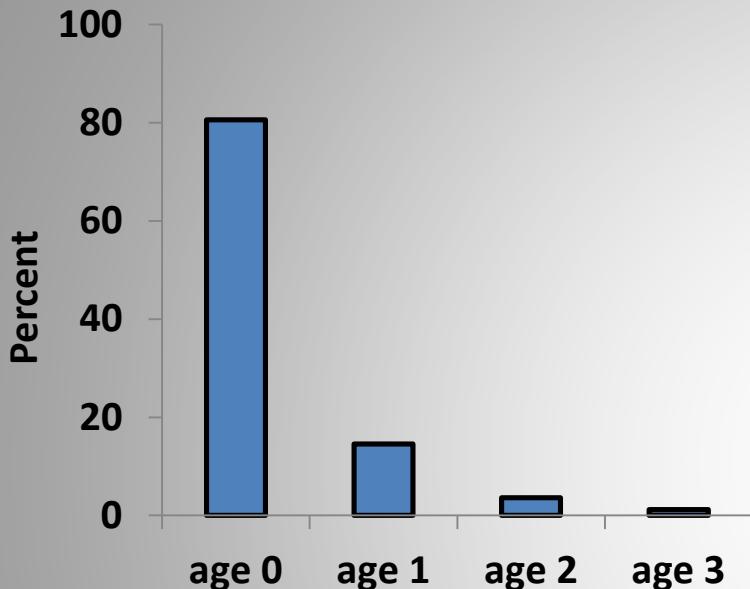
BUT, 50% parr progeny were **AxA** cross



Survival 1.3%

## Results – Successful phenotypes?

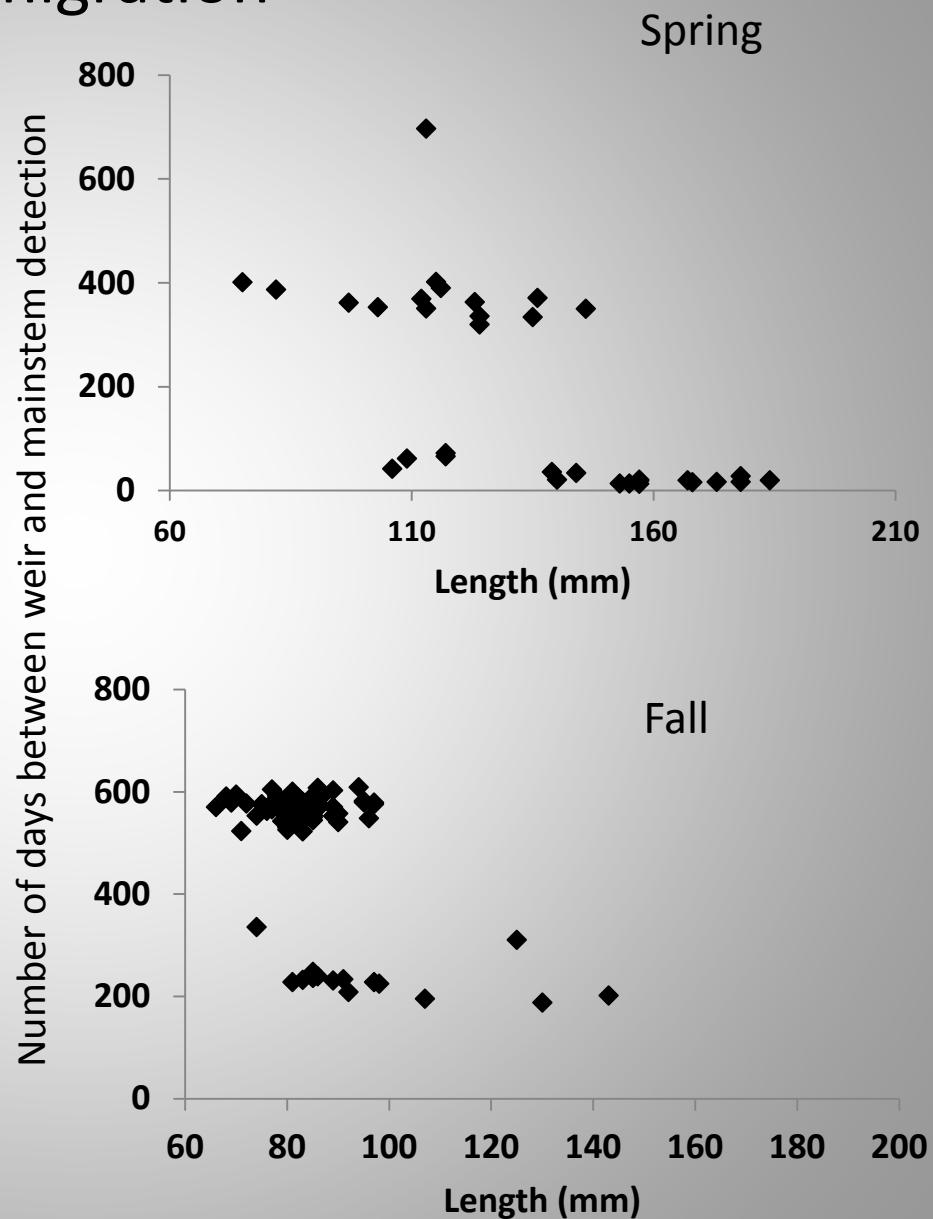
### Juvenile migration



80% offspring captured

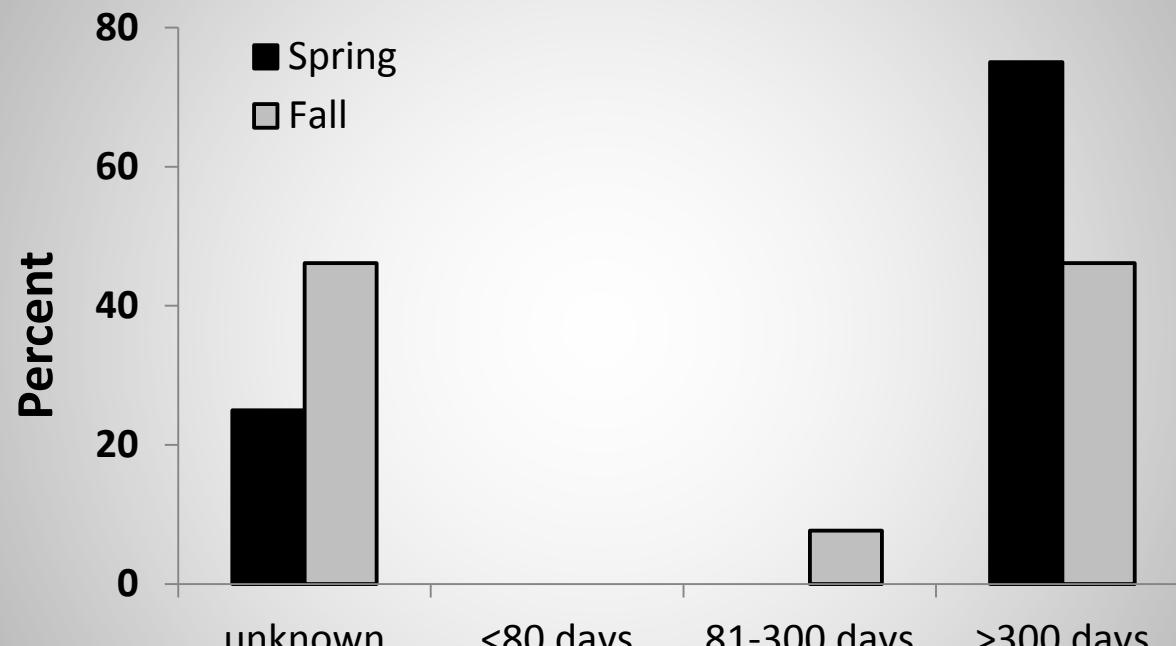
Age 0

Most parr reared >200 days downstream from the natal tributary before smolt outmigration



Results – Successful phenotypes?  
**Juvenile Migration/Survival**

Most successful **adult returns** rear in –basin > 300 days

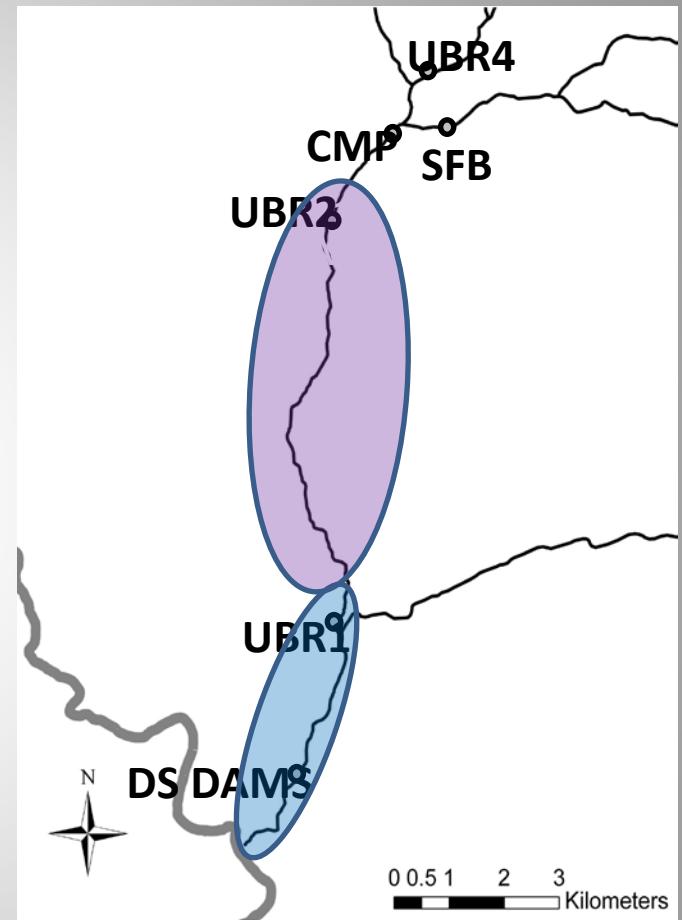


Days between BC outmigration and smolt detection

## Results – Spatial extent of measurable changes?

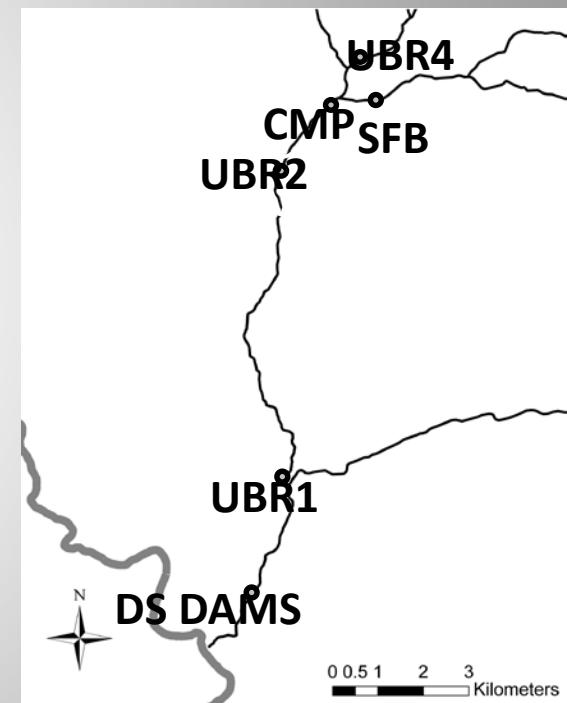
Site	Before Year	After Year	$F_{ST}$	Pval
DS Dam	2005	2009	0.014*	0.001*
UBR1	2004	2008	0.021*	<0.001*
UBR1	2004	2009	0.027*	<0.001*
CMP	2005	2009	0.002	0.047
UBR4	2004	2008	0.011*	0.009*
UBR4	2004	2009	-0.002	0.558
SFB	2005	2008	0.004	0.121
SFB	2005	2009	0.002	0.276

\* Indicates statistical significance

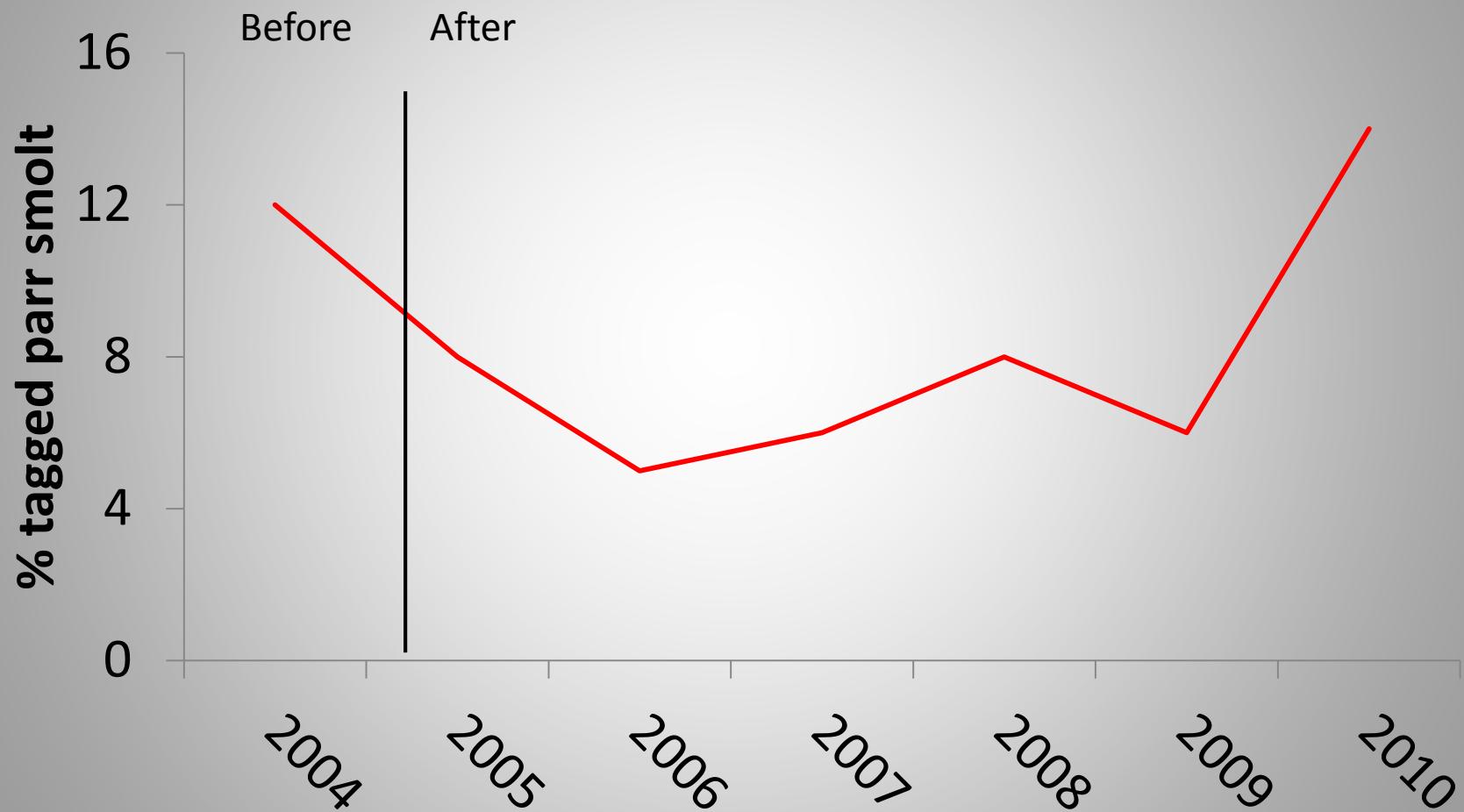


## Results – Temporal Tests (sampling effect)

site	year	year	$F_{ST}$	Pval
UBR1	2008	2009	-0.003	0.253
UBR2	2008	2009	-0.004	0.880
UBR4	2008	2009	<-0.001	0.147
SFB	2008	2009	0.005	0.568



# % smolt UBR1



# Conclusions - Colonization



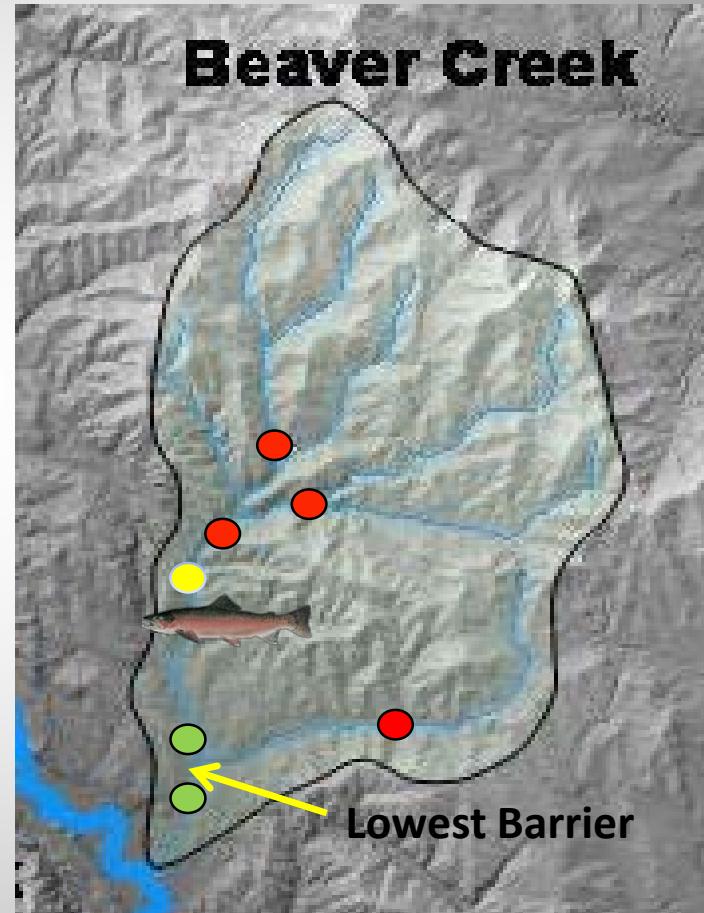
## Before barrier removal

- Fragmented
- Smolts out, but no adult returns



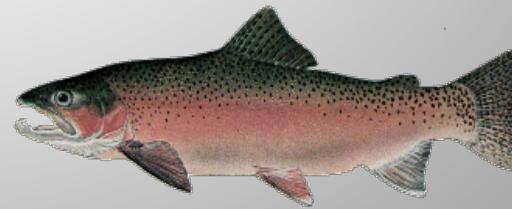
## After barrier removal

- Genetic shifts lowest site km 5
- Tag movement to middle site km 12
- Smolts out and adult returns



# Conclusions – Indiv Success

- Fluvial RBT were key to re-colonization, gene flow and reserve of wild genotype
- Hatchery unsuccessful in early years of colonization
- Phenotypes of successful spawners shifted dramatically between 2005 and 2006
- Successful juvenile SH reared in the Methow R (or Wells Res) for 1 to 2 years prior to smolt outmigration
- No clear relationship between number of offspring and returning adults



# What is the **weakest link** in steelhead conservation?

- Phenotypes/genotypes under selection for adult survival and how these interact in time, space, density -> selection gradients
  - Interactions between rearing habitats and selection gradients influences fitness
- Aggressive work to understand, control or eliminate hatchery steelhead effects



# Acknowledgements

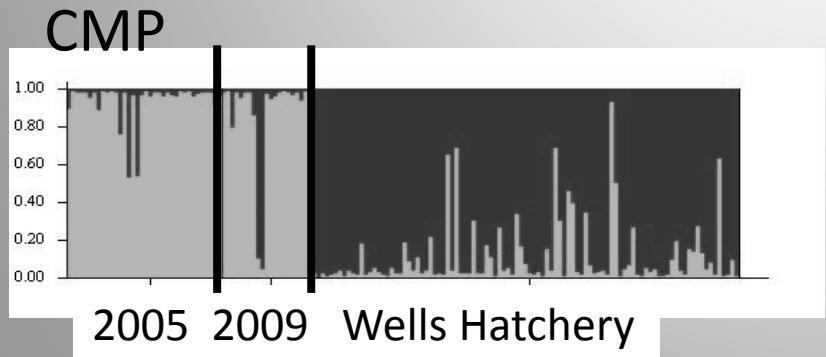
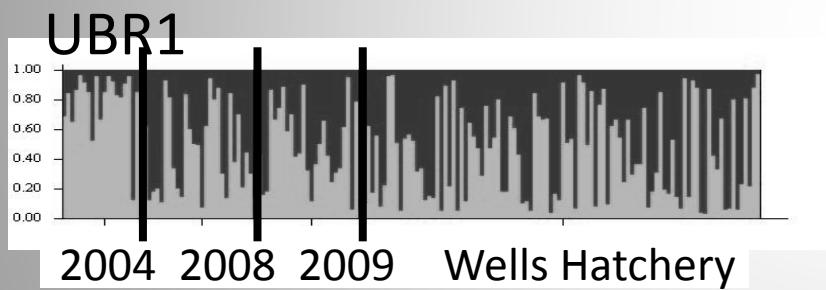
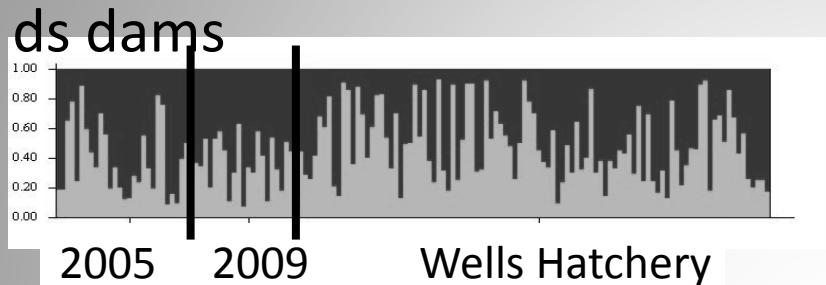
Funding provided by the Bureau of Reclamation

Data collection assistance (USGS) –  
Kyle Martens, Wes Tibits, Brian Fisher

Lab analysis assistance Joyce Faler (Univ. of Idaho)

Coordination Greg Knott, Michael Newsom

# Results - % Hatchery



Site	Before %	After %
DS Dams	40	35
UBR1	27	47
CMP	6	12

# Gene Flow



## Genetic Differentiation (Fst)

	Fluvial	Wild Anad
Wild Anad	0.002 - ns	
Hatchery	0.006	0.004