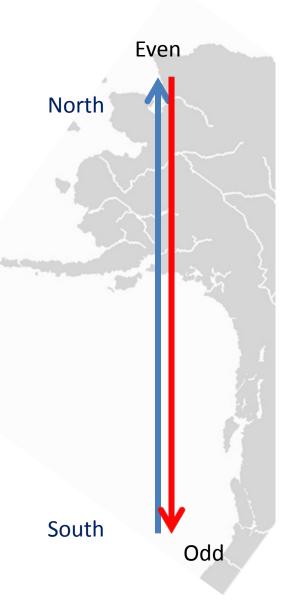


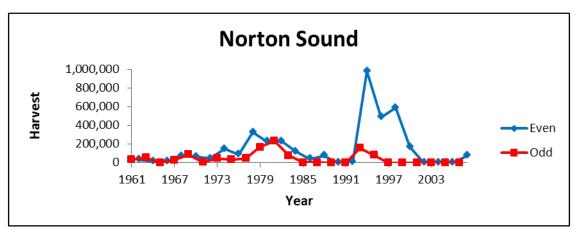
### What's Unique About Pink Salmon?

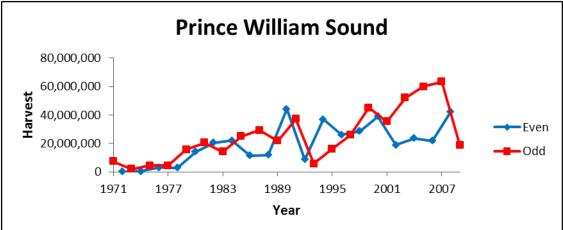
- Strict two-year life cycle-no gene flow
- Dominance of one broodline in many areas
- Currently high abundance, colonizing

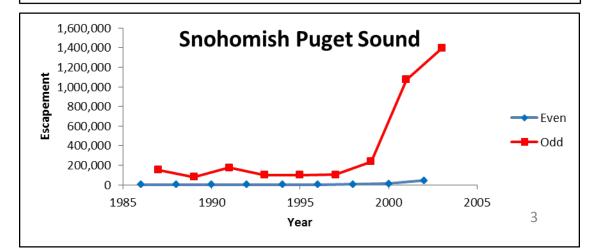


#### **Broodline Dominance**

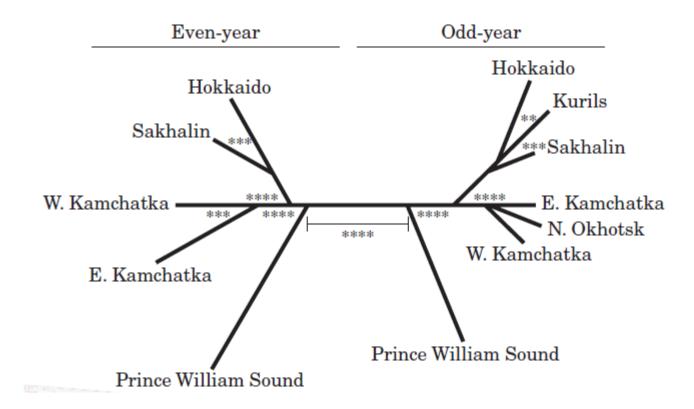






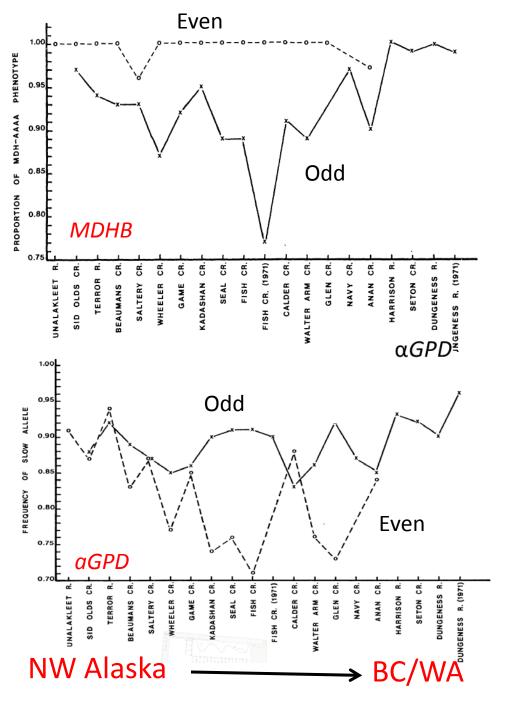


#### ODD- AND EVEN-BROODLINE ASIAN PINK SALMON



Hawkins et al. (2002)





- Aspinwall (1974)
  - allele frequency differences between even and odd in similar locations supports neutral-random drift hypothesis rather than balancing selection

#### What Can Genomics Add?

- Signals of adaptation (selection)
- Can broodlines be used as replicates to identify adaptive loci?
- Provide higher resolution markers



### Paired even/odd collections

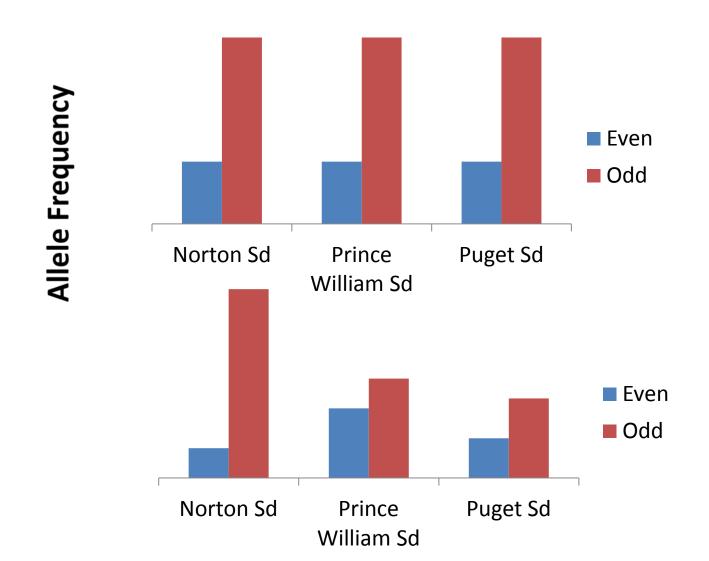
Nome R, Norton Sound 1994/1991

- Geographically dispersed pairs of even/odd populations
- Temporal replicates
  - Sample sizes 20 to 24
- What are expectations for neutral and selective markers?

Koppen Cr, Prince William Sound 1996/1991

> Snohomish R, Puget Sound 1996/2003

## Expectations of Neutral SNP Reflect Broodline and Geographic Divergence



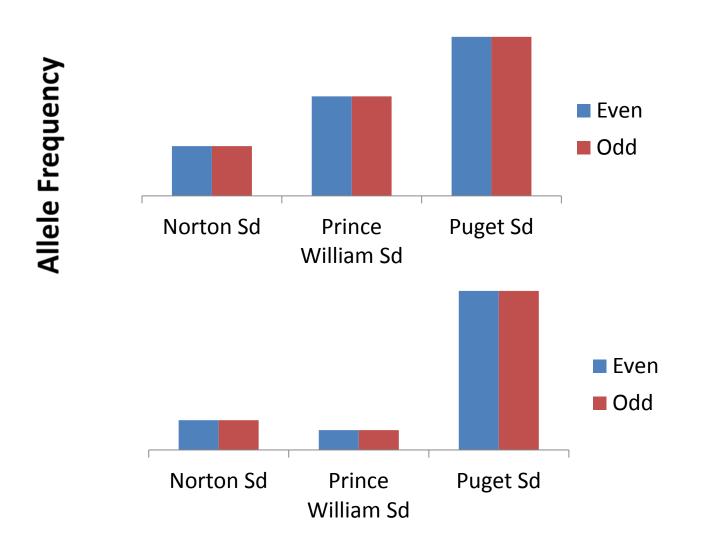
### Share local environmental conditions, pathogens exposure



#### Share spawning location, timing, substrate



## Expectation of SNPs Reflecting Local Adaptation & Selection



### Pink RAD Pipeline

#### Restriction-site Associated DNA

- Identified over 70,903 putative SNPs
  - 1 bp differences, exclude close variants (PSVs)
- Assign genotypes from sequence counts
- Final data set after conservative filtering

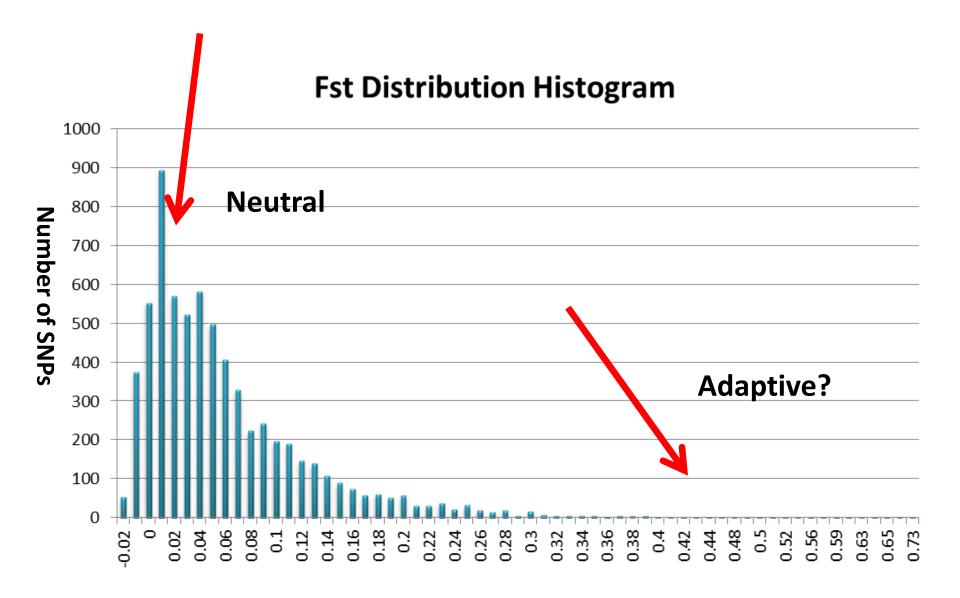
131 of 140 Individuals 6,743 Loci

```
AGCTGACTTGACCT C GAAGTGCATTGCTAGGTCGCGATTG
AGCTGACTTGACCT A GAAGTGCATTGCTAGGTCGCGATTG
```

AGCTGACTTGACCT C GAAGTGCATTGCTAGGTCGCGATTG
AGCTGACTTGACCT A GAAGTGCATTGCTAGGTCGCGATTG

A:C Heterozygote



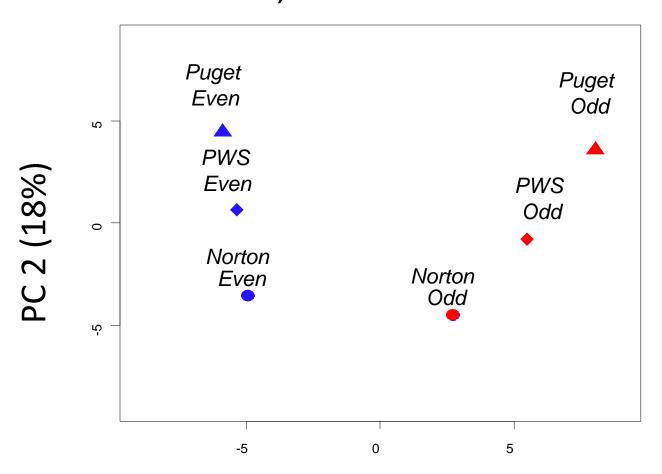


### Diversity Among North American Pink Salmon 6,743 SNPs

Broodline	Fst
Even	0.036
Odd	0.053
Even + Odd	0.086

Pairwise by Location	Fst
Norton Sd	0.065
Prince William Sd	0.098
Puget Sd	0.146

#### Pink Salmon PCA 6,743 SNPs



PC 1 (52%)

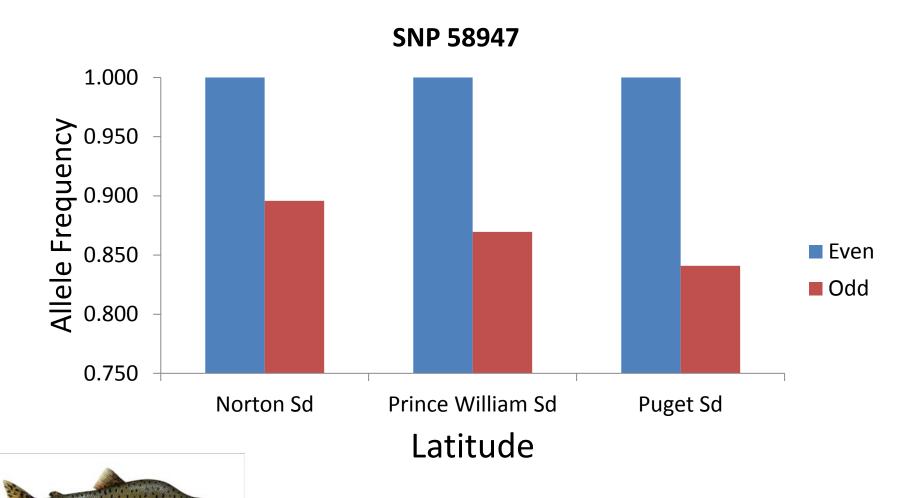
### Eignenvector Loadings Correlation Coefficients

- PC 1
  - Explains 52% of variance
  - 244 significant SNPs



Critical Value (0.95)

## Eigenvector Correlations with PC 1 Reflecting Geographic and Broodline



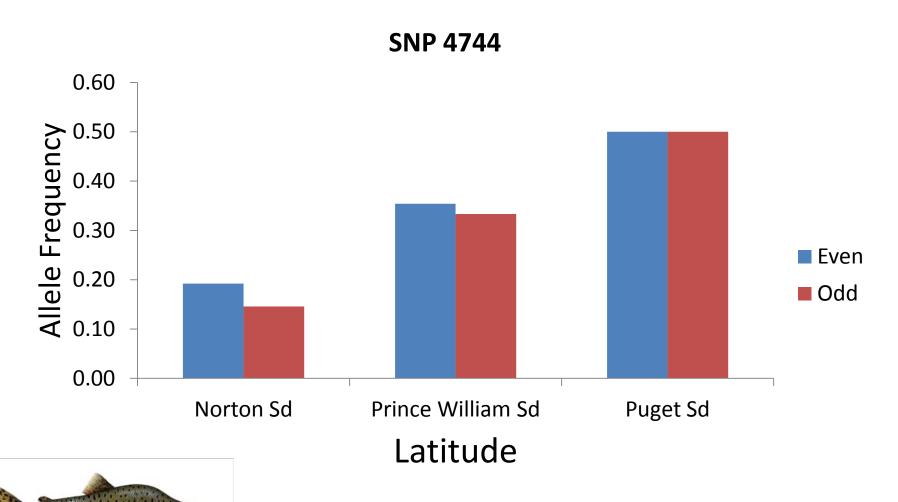
### Eignenvector Loadings Correlation Coefficients

- PC 1
  - Explains 52% of variance
  - 244 significant SNPs
- PC 2
  - Explains 18% of variance
  - 27 significant SNPs



Critical Value (0.95)

# Eigenvector Correlations with PC 2 SNP Reflecting Local Adaptation?



#### Worth Another Look--Yes

- Identified 6,743 candidate SNPs
  - 27 with potential functional significance
    - Adaptive Markers, high GSI potential
- More diversity in odd
  - Even/odd more divergent in south
- Adaptive differences between broodlines?
  - One year derived from other?
  - Independent colonization and expansion?



#### Pink Takeover in Pacific Northwest?

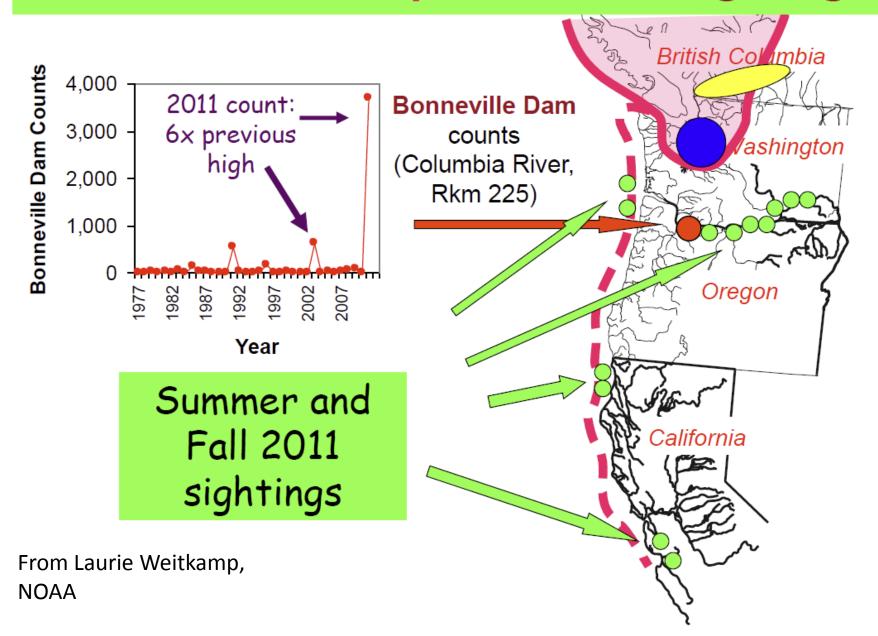
- Dramatic increase in last decade in Puget Sound and Columbia River in odd-year
- Abundant recreational opportunities
- Concerns that they may negatively affect ESA-listed populations





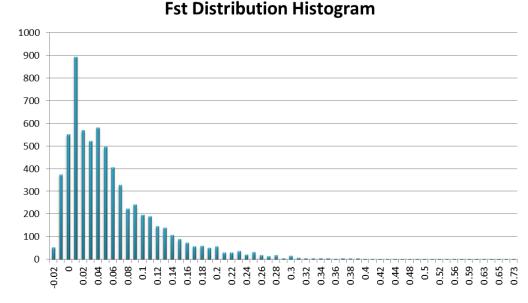
Puget Sound, Washington

#### Aside: 2011 adult pink salmon sightings



#### What's next?

- High-throughput assays for genotyping
  - High conversion rates directly from RAD data (approx. 90% to TaqMan)
- UW, WDFW, and NOAA (NWFSC) collaborating on SNP panels and baseline for Pacific Northwest



### **Potential Applications**

- Track expansion and colonization
  - Origin of colonizers
- Genetic Mark Recapture hatchery/wild interactions
  - Investigate straying, reproductive success
- Migration and stock interactions on the high seas
  - Expand range-wide panel
  - PacSNP Pink

