

Status of Agency GSI Databases and recommendations

Habicht/Beacham/Moran/Narum
(Genetics)

Sockeye salmon loci in DFO baseline

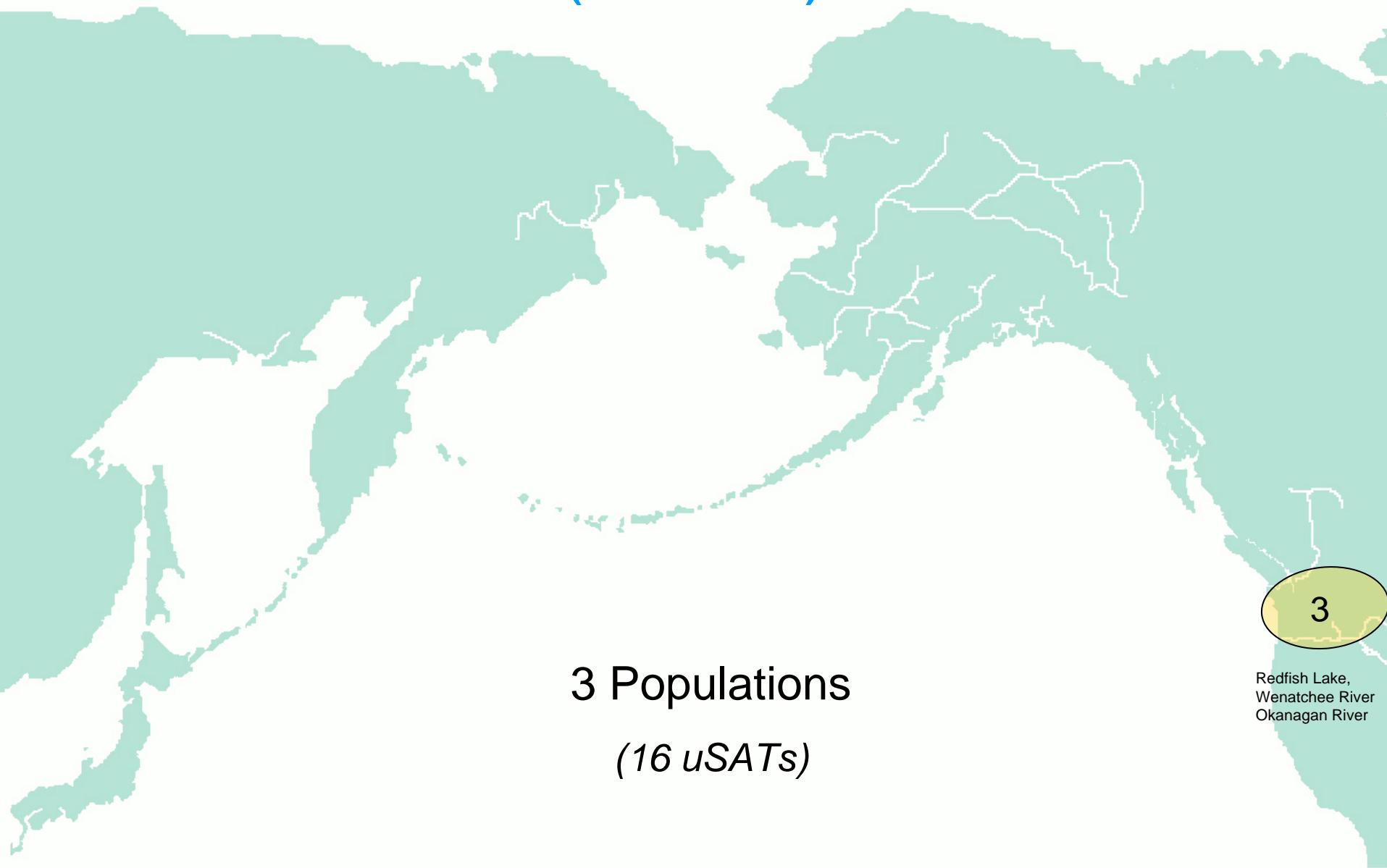
| Locus | Alleles |
|----------------|---------|
| <i>Oki1a</i> | 8 |
| <i>Oki1b</i> | 10 |
| <i>Ots107</i> | 15 |
| <i>Omy77</i> | 20 |
| <i>Ots2</i> | 26 |
| <i>Ots3</i> | 26 |
| <i>Oki16</i> | 26 |
| <i>Ots108</i> | 29 |
| <i>Ots103</i> | 30 |
| <i>One8</i> | 32 |
| <i>Ots100</i> | 33 |
| <i>Oki6</i> | 37 |
| <i>Oki29</i> | 39 |
| <i>Oki10</i> | 83 |
| <i>DAB -β1</i> | 15 |

Sockeye salmon: Summary of the number of sampling sites or populations within geographic regions from Beacham et al. (2006). Range of annual and population samples sizes within regions is in parentheses. Fourteen microsatellite loci and an MHC locus were surveyed.

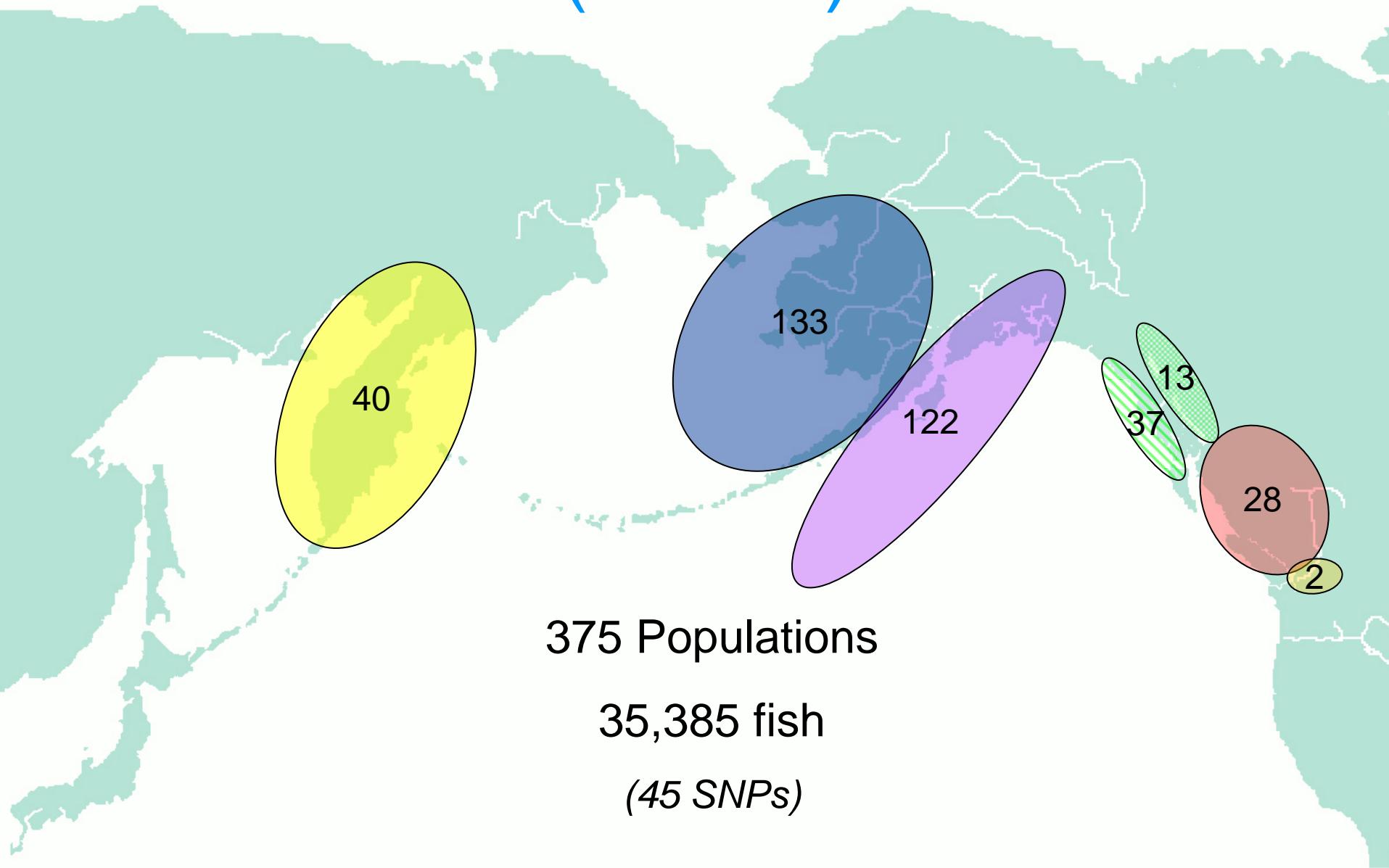
| Region | Number of populations | Mean annual sample size | Mean population |
|----------------------|--------------------------|-------------------------|-----------------|
| | | | sample size |
| Columbia River | 2 | 71 (15, 194) | 285 (68,502) |
| Washington | 3 | 114 (50, 201) | 114 (50, 201) |
| Fraser River | 53 | 94 (5, 400) | 270 (15, 858) |
| West coast Vancouver | 15 | 90 (19, 197) | 132 (19, 279) |
| Island | | | |
| Nimpkish River | 3 | 108 (42, 290) | 288 (203, 367) |
| Southern BC | 6 | 114 (12, 219) | 171 (18, 325) |

| | | | |
|------------------|----|---------------|----------------|
| Central BC | 16 | 79 (27, 223) | 97 (27, 223) |
| Owikeno Lake | 10 | 77 (7, 114) | 224 (86, 398) |
| Long Lake | 3 | 99 (39, 205) | 297 (139, 490) |
| Queen Charlotte | 5 | 71 (41, 99) | 114 (41, 190) |
| Islands | | | |
| Nass River | 11 | 96 (24, 264) | 313 (40, 797) |
| Skeena River | 14 | 78 (33, 200) | 151 (33, 287) |
| Babine Lake | 11 | 95 (54, 200) | 208 (78, 499) |
| Unuk River | 1 | 50 (50,50) | 50 (50,50) |
| Stikine River | 17 | 83 (6, 405) | 152 (26, 474) |
| Taku River | 10 | 57 (12, 100) | 86 (12, 199) |
| Alsek River | 15 | 83 (10, 238) | 144 (10, 592) |
| Southeast Alaska | 20 | 151 (45, 343) | 197 (45, 300) |

2007 Sockeye Microsatellite Baseline (NOAA)



2007 Sockeye SNP Baseline (ADFG)



Coho salmon loci in DFO baseline

Locus

Ots2

Ots3

Ots101

Ots103

Oki1

Oki10

Oki100

Oki101

Ogo2

Omy325

One111

OtsG253b

Ssa408

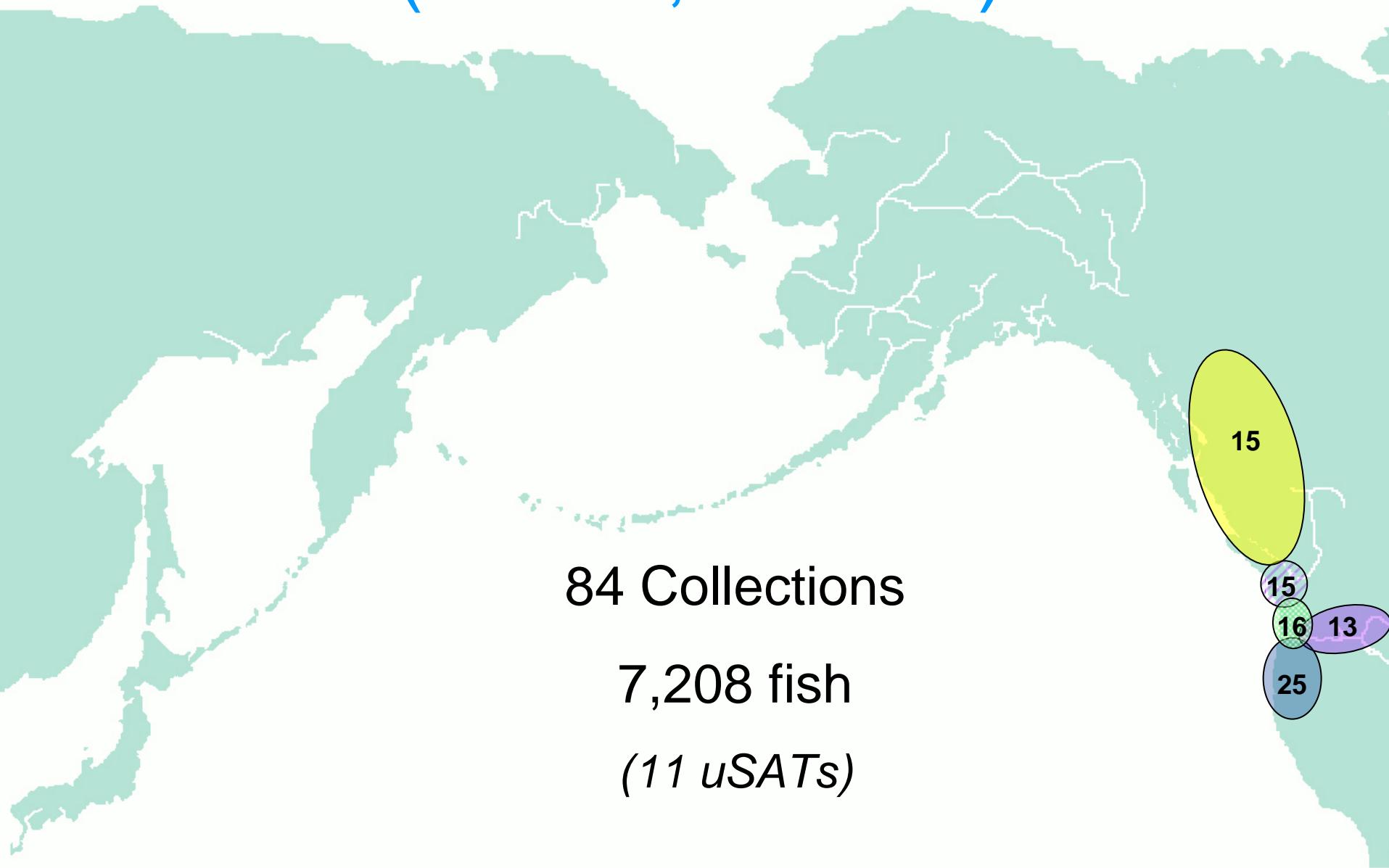
Alpha1

Alpha2

Coho salmon: Summary of the number of regions, sampling sites or populations within geographic regions, and individuals surveyed. Thirteen microsatellite loci and two MHC loci were surveyed.

| Region | # of collections | # of coho |
|---------------|------------------|-----------|
| Transboundary | 7 | 700 |
| SE Alaska | 9 | 1450 |
| QCI | 20 | 1400 |
| Nass | 3 | 750 |
| Skeena | 29 | 4500 |
| Central Coast | 40 | 5850 |
| South Coast | 28 | 3650 |
| ECVI | 22 | 6350 |
| WCVI | 13 | 4300 |
| Fraser | 47 | 13,300 |
| Washington | 15 | 1400 |
| Columbia | 9 | 800 |
| Oregon | 10 | 850 |
| California | 4 | 150 |

2007 Coho Microsatellite Baseline (NOAA, Seattle)



| Locus | NMFS | NMFS | CDFO/WDFW | USFWS | USFWS | OSU | BM | Allele ladder Candidates |
|--------------|------------|------------|---------------|-----------|--------|-----|----|--------------------------|
| | Manchester | Santa Cruz | collaboration | Abernathy | Alaska | | | |
| Ocl8 | X | X | X | X | | | | 1 |
| Oki1 | X | X | X | X | X | | X | 1 |
| Oki10 | X | | X | X | | | | 2 |
| Oki23 | X | | | X | | | | 3 |
| One13 | X | X | | X | | X | X | 2 |
| Ots103 | X | X | X | X | | X | X | 1 |
| Ots213 | X | | X | X | | X | | 2 |
| Ots3 | X | | | X | | s | X | 3 |
| OtsB3 | X | | | X | | | | 3 |
| OtsG422 | X | X | | X | | | | 2 |
| P53 | X | | X | X | | X | X | 1 |
| iso-Ots2 | s | | | | X | | | |
| Oki11 | s | | | | | | | |
| Oki13 | s | X | | | | | | |
| Oki2 | s | | | | | | | |
| Oki3 | s | | | | X | | | |
| Ots101 | s | | X | | | | | 3 |
| Ots105 | s | X | | | X | | | 3 |
| Ots2 | s | | | | | X | X | |
| Ots208 | s | | | | | | | |
| Ots212 | s | | | | | s | | |
| OtsG249 | s | | | | | | | |
| OtsG253b | s | | X | | | | | 3 |
| OtsG3 | s | X | | | | | | |
| OtsG68 | s | X | | | | | | |
| OtsG78b | s | X | | | | | | |
| OtsG83b | s | X | | | | | | |
| Ogo1a | | | | | | | | |
| Ogo2 | | | | | | | | 3 |
| Oke2 | | | | | X | | | |
| Oke3 | | | | | X | | | |
| Oke4 | | | | | X | | | |
| Oki100 | | | X | | | | | 3 |
| Oki101 | | | X | | | | | 3 |
| Oki16 | | | | | | X | | |
| Omm1121 | | | | | | | | |
| Omm1128 | | | | | | | | |
| Omy1011 | | | X | | | | | 3 |
| Omy116 | | X | | | | | | |
| Omy325 | | | X | | | | | 3 |
| Omy77 | | | | | | | | |
| One111 | | | X | | | | | 3 |
| One11b | | X | | | | | | |
| One13M | | | X | | | | | 3 |
| One2 | | | | | | | | |
| One3 | | | | | X | | | |
| Ots1 | | | | | | s | | |
| Ots10 | | | | | | s | | |
| Ots108 | | X | | | | | | |
| Ots1b | | X | | | | | | |
| Ots206 | | | | | | | s | |
| Ots208b | | | | | | | s | |
| Ots209 | | | | | | | s | |
| Ots215 | | | | | | | X | |
| Ots2M | | | X | | | | | 3 |
| Ots3.1 | | | X | | X | | | 3 |
| Ots3M | | | | | | | | |
| Ots9 | | | | | | | | |
| Ssa14 | | X | | | | | | |
| Ssa407 | | | X | | | | | 3 |
| Ssa85 | | X | | | | | | |
| Total in use | 11 | 17 | 18 | 11 | 9 | 8 | 7 | |

X = locus is in use

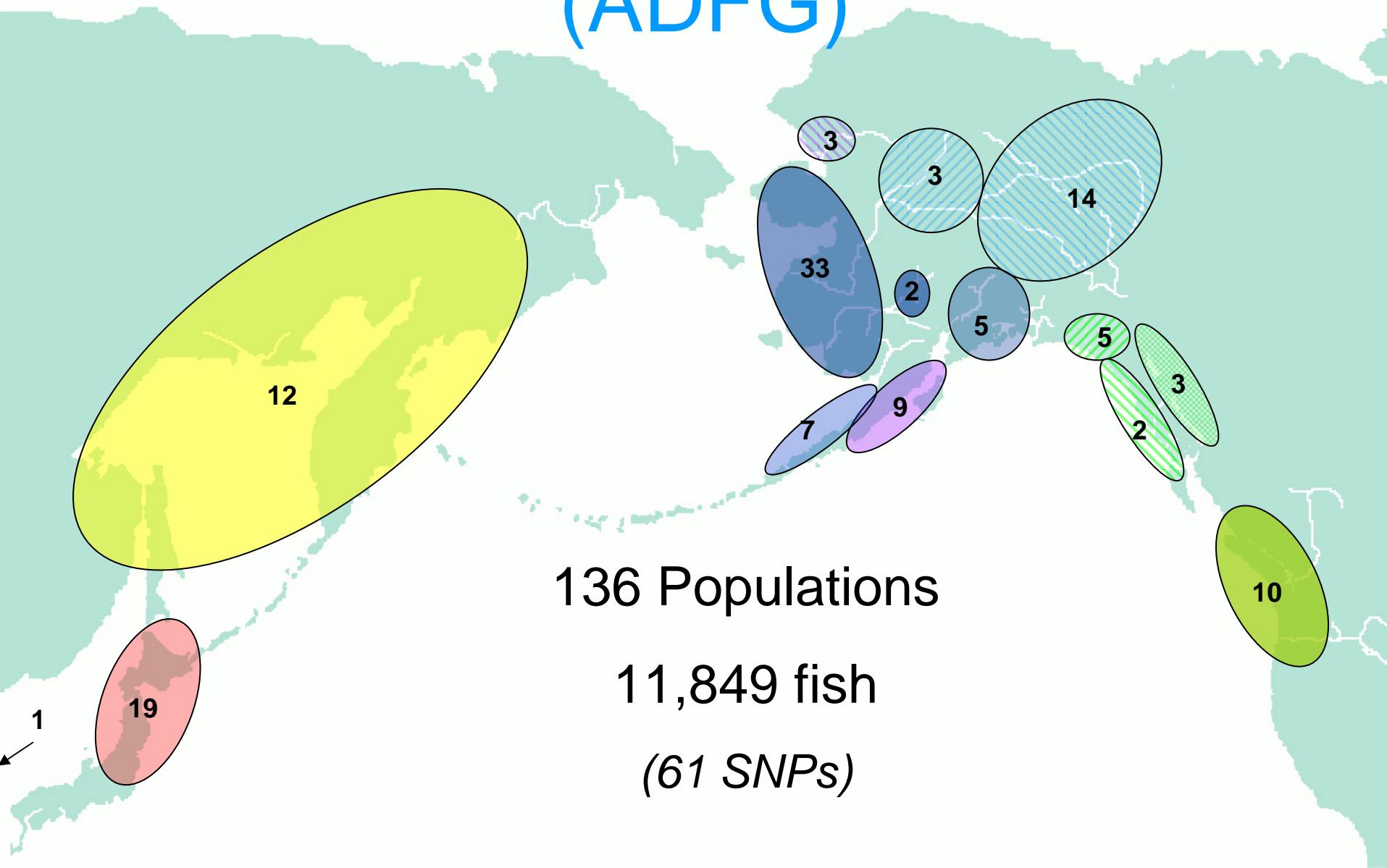
s = locus has been screened and is being or has been evaluated for possible use

Coho salmon uSATs – overlap among laboratories

2007 Coho SNP Baseline (ADFG)



2007 Chum SNP Baseline (ADFG)



GAPS Chinook

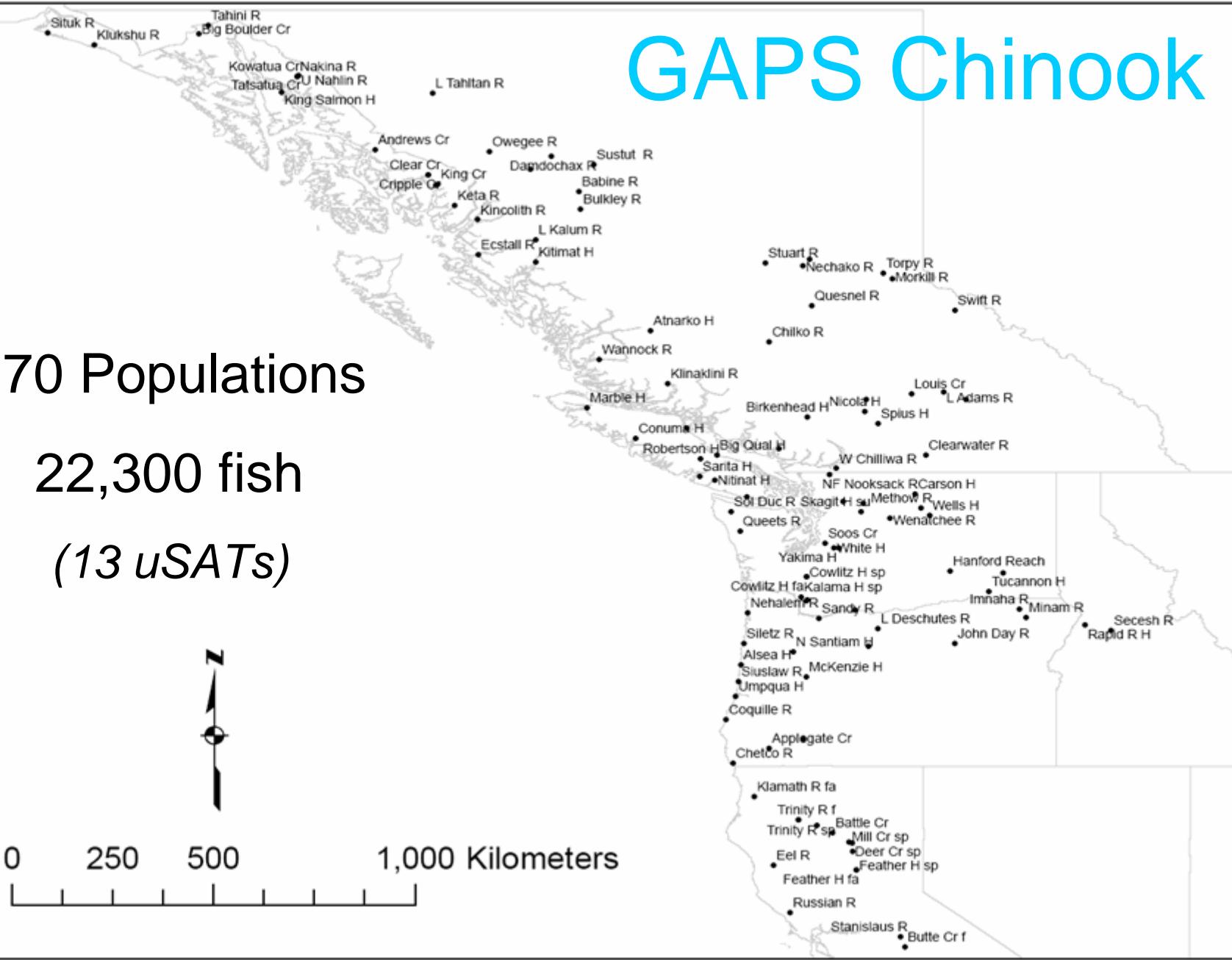
170 Populations

22,300 fish

(13 uSATs)



0 250 500 1,000 Kilometers

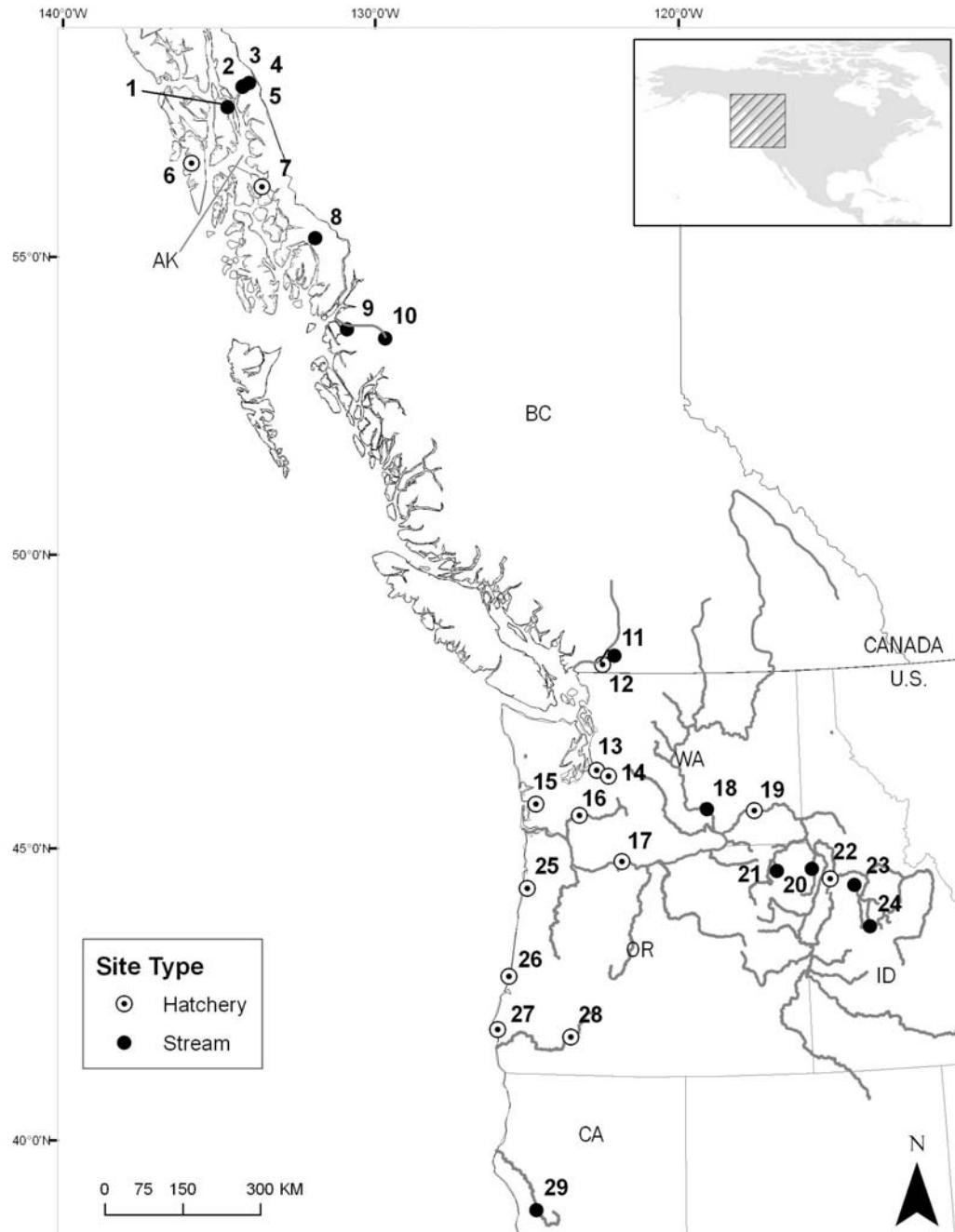
A horizontal scale bar with tick marks at 0, 250, 500, and 1,000 kilometers. The first two segments each represent 250 km, and the third segment represents 500 km, totaling 1,000 km.

2007 Chinook SNP Baseline

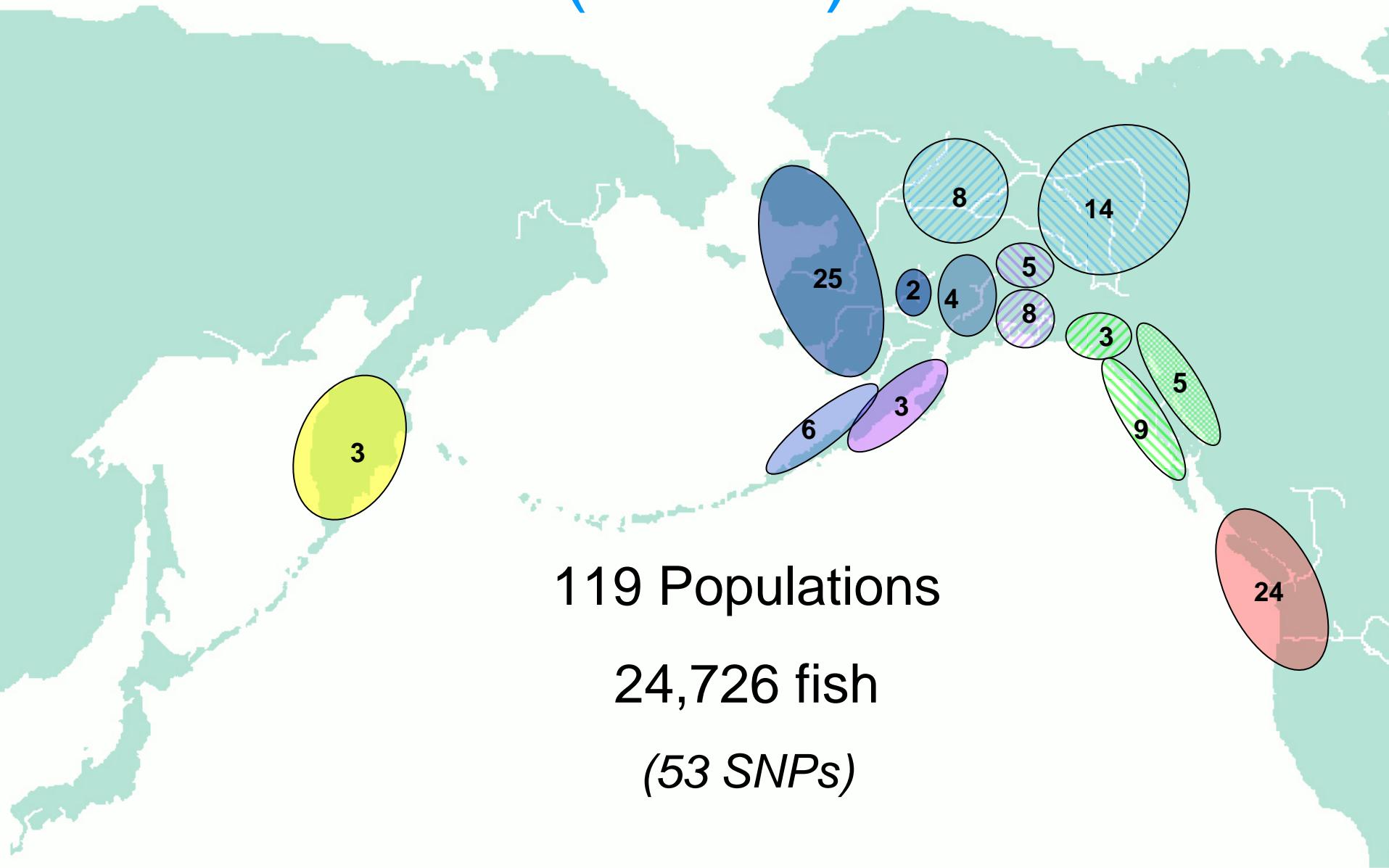
Narum et al.

29 Populations

2,782 fish
(37 SNPs)



2007 Chinook SNP Baseline (ADFG)



SNP assays available for Pacific Salmon

| Species | Number of genotyping assays |
|----------------|-----------------------------|
| Chinook salmon | 70+ |
| Coho salmon | 19 |
| Chum salmon | 77 |
| Sockeye salmon | 44 |
| Pink salmon | 0 |

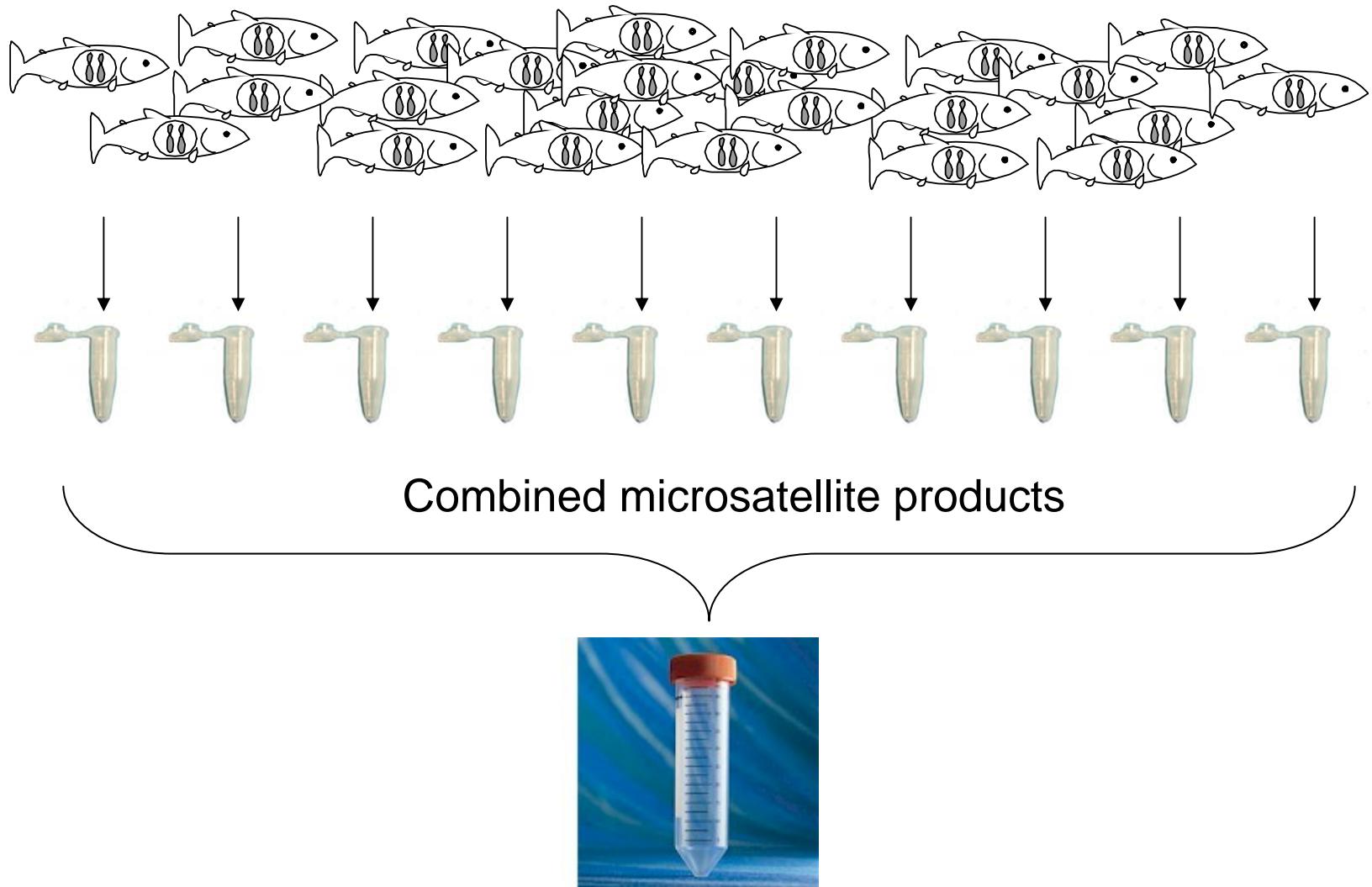
Conclusions/Recommendations

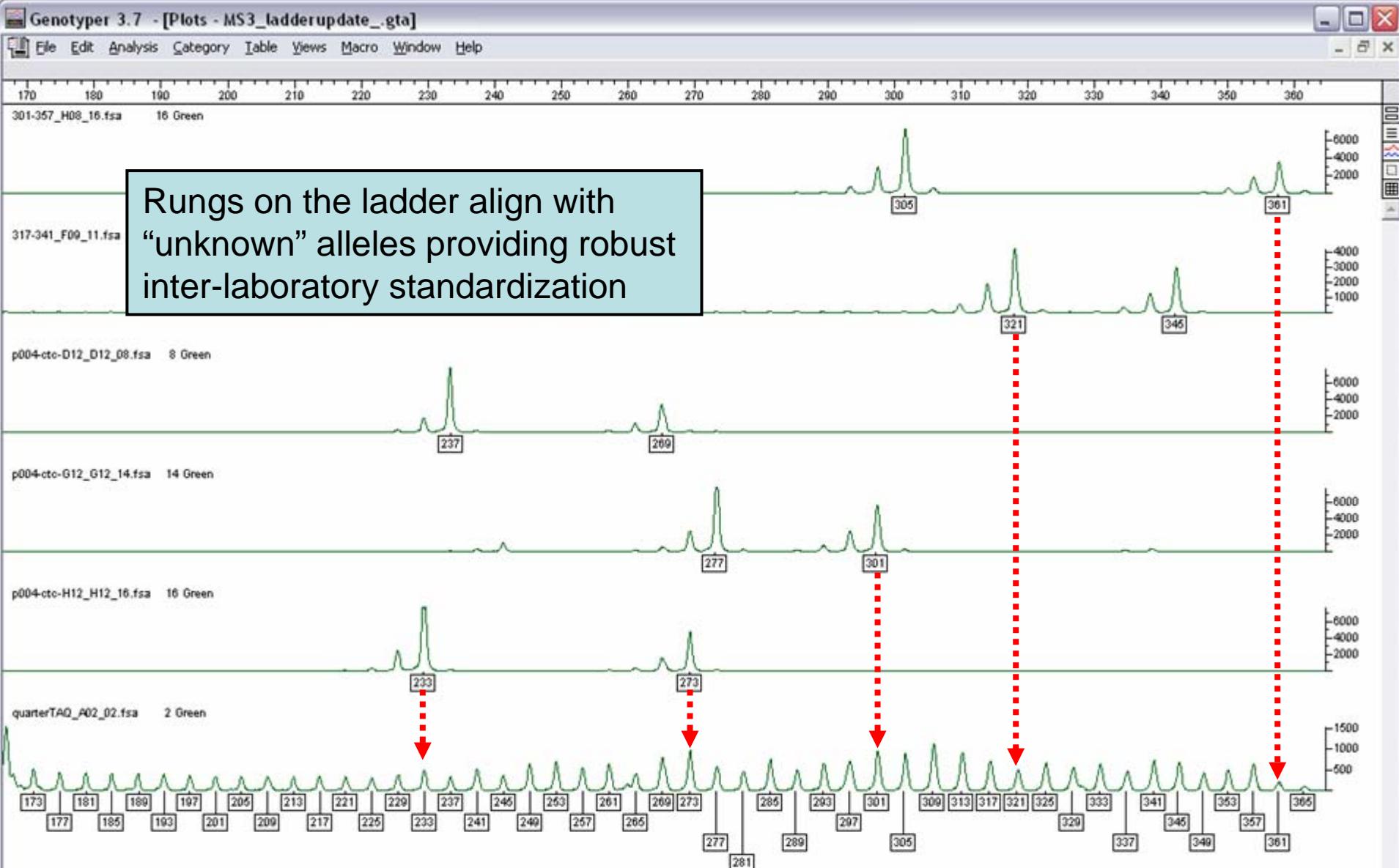
- Both SNP and microsatellite databases are extensive for sockeye, chinook, and chum salmon.
- Empirically test cost-benefit of markers both regionally and coastwide.
- Standardize loci and allele scoring.
- Christian Smith will provide group's recommendations on marker choice.

Standardizing coho data

- Current proposal to the Southern Fund to standardize existing coho microsatellite data
- Consolidate standardized and non-standardized coho data in the existing GAPS database
- Allele ladders will dramatically simplify microsatellite standardization and certification of new labs

What is an Allele Ladder?





Please see LaHood et al. 2002 (Molecular Ecology Notes 2:187–190) for use of ladders in inter-laboratory standardization