

Stock Identification of Chinook Salmon from Southeast Alaska: A review of the past and a look forward

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Alaska Department of Fish and Game



SE Alaska Troll Fishery

1999-2006

Fishing Seasons:

October-December

January-April 14

April 15-June

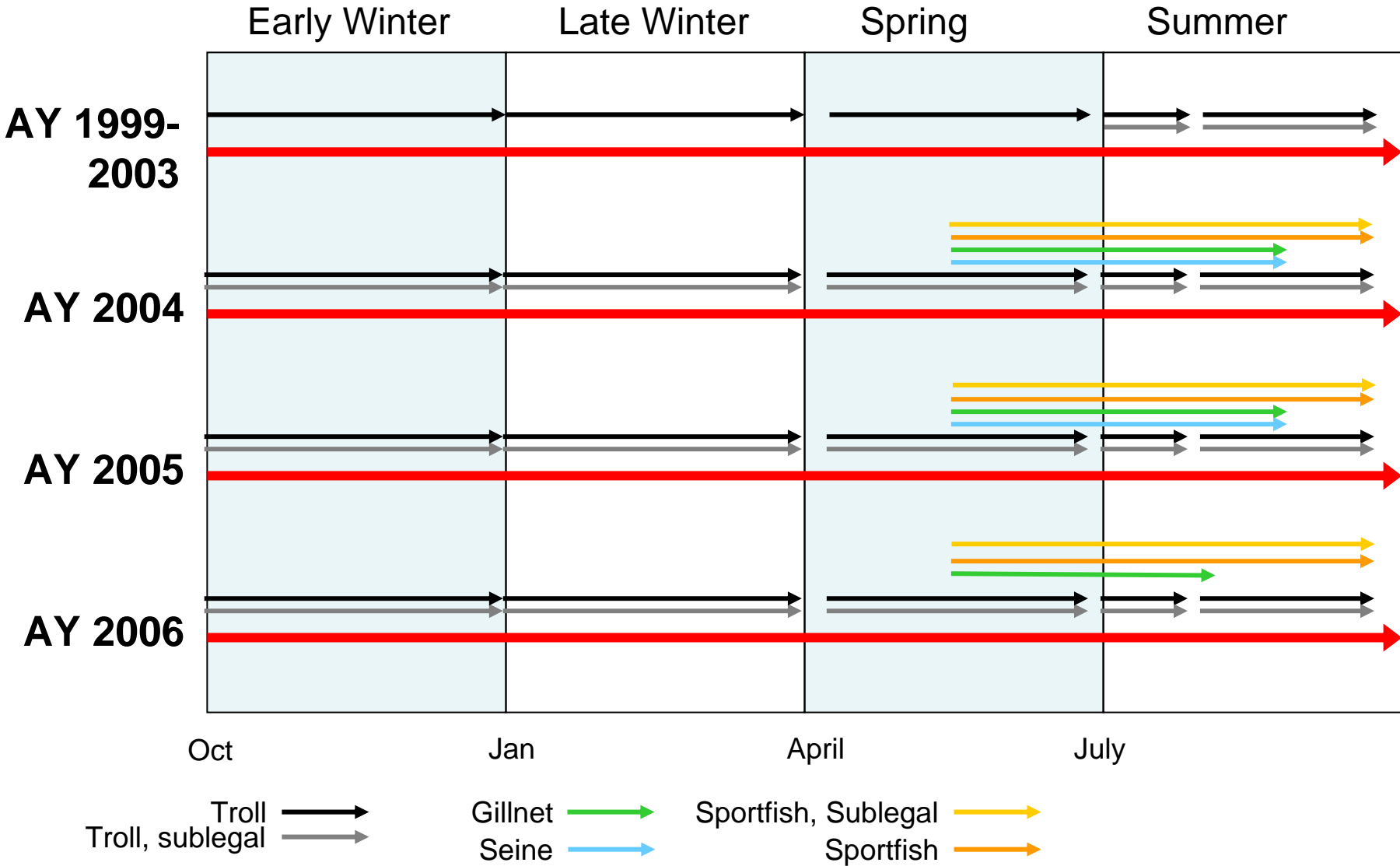
July-September

Legal-sized Chinook (>28 in)

Sublegal Chinook (<28 in)

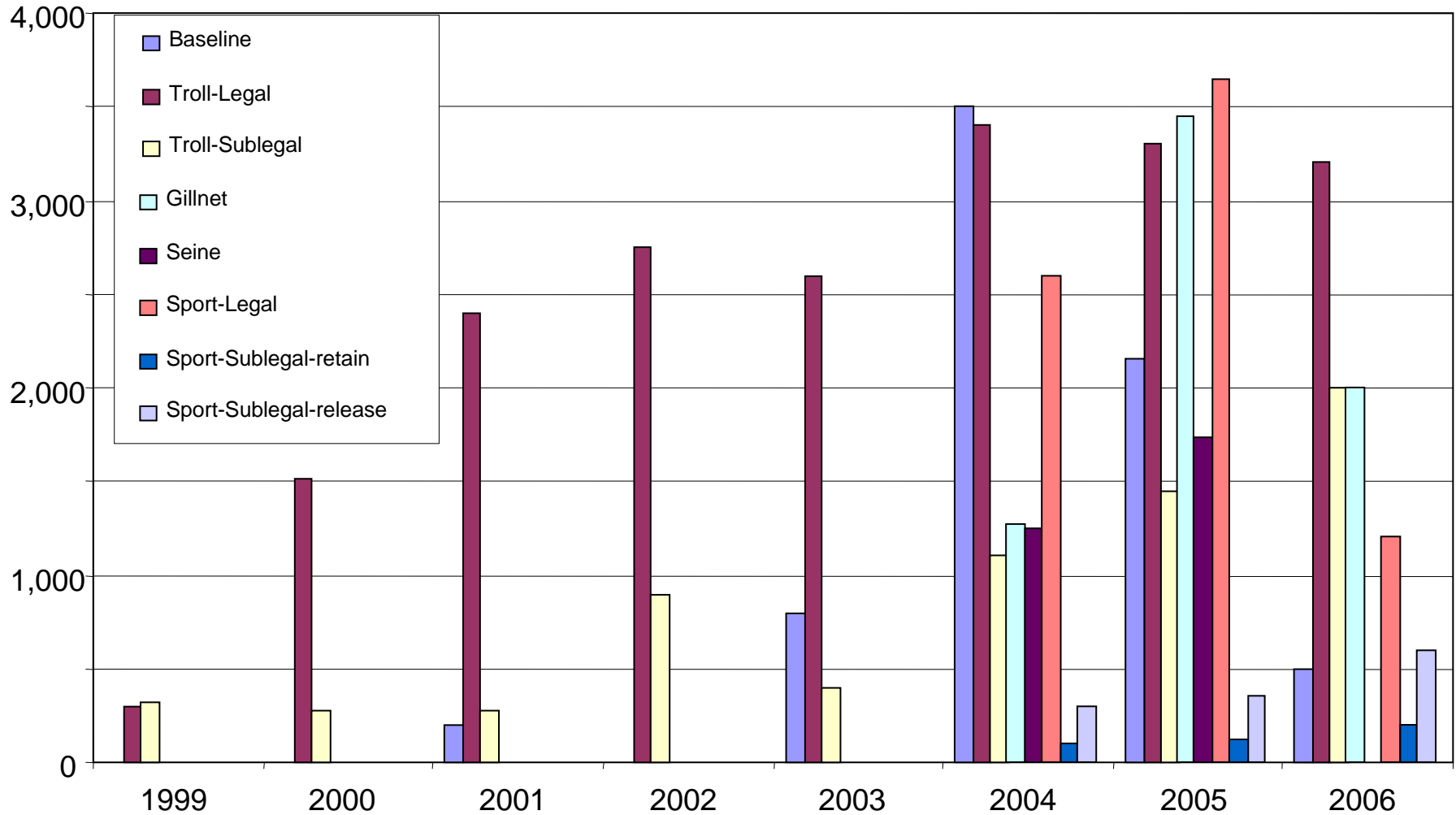


Sampling of SE Alaska Chinook Fisheries



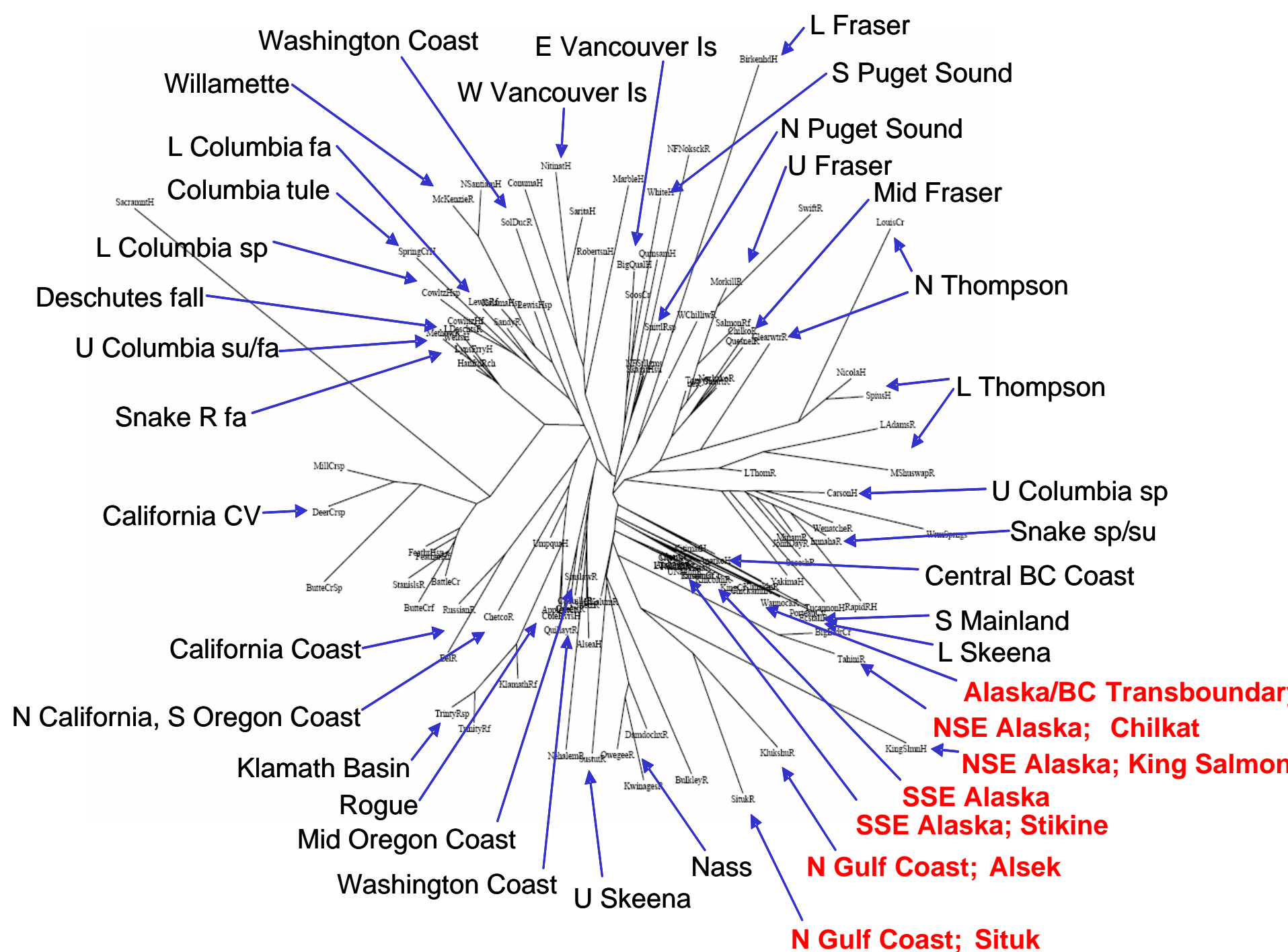
SE Alaska Chinook sampling effort

1999 - 2006



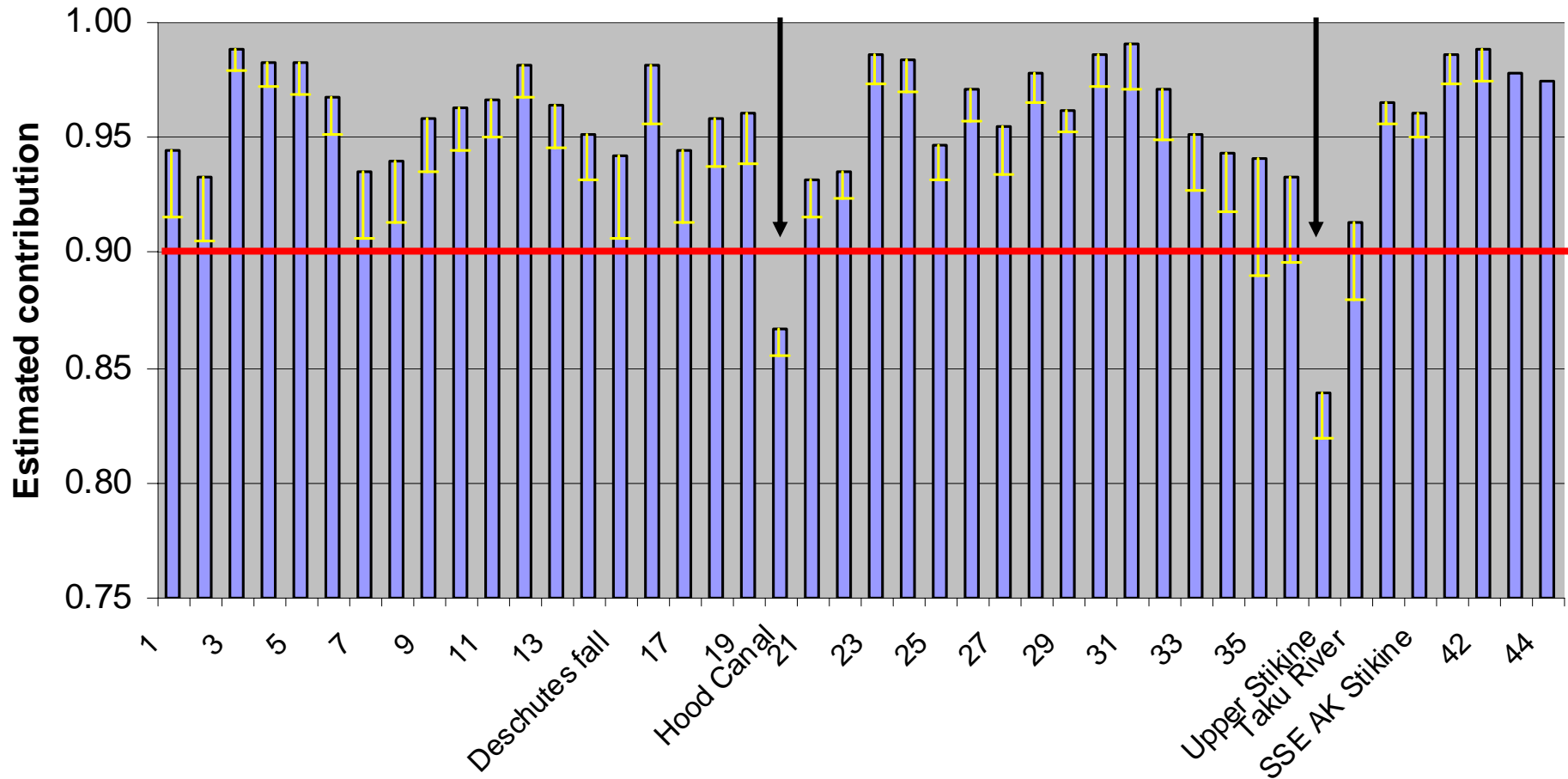
What are the questions?

- Composition of catch spatially, temporally within and among years:
 - Troll
 - Sport
 - Gillnet, seine
- Sublegal vs. legal sized



100% sim results (SPAM)

GAPS version 2.1



Options:

- 1) Add more individuals and/or populations
- 2) Add more genetic markers
- 3) Combine with a genetically similar group

Challenges

Study Design

- Sampling difficulties
 - Coordinating proportional sampling in diverse and remote ports across SEAK
- Improvement of markers
 - 1999 – 2003, Allozymes
 - 2004 – present, Microsatellites

Troll Legal Sampling Design

2007

Port	Early Winter October 11 – Dec. 31	Late Winter January 1 – April 14	Spring May 1 – June 30	Summer July 1 – September 30 1 st ret./ 2 nd ret.
Sitka	400	350	300	300/300
Yakutat	30	30		30/30
Juneau	30	30	200	
Port Alexander				60/60
Elfin Cove				30/30
Pelican				30/30
Hoonah			75	40/40
Wrangell			300	
Petersburg	25	40	100	60/60
Ketchikan	40	60	200	100/50
Craig	20	20		150/50
Projected totals	545	530	1175	800/650
Overall total	3975			

Troll Sampling Design

9 Ports

Size Class	Early Winter Oct 11 – Dec. 31	Late Winter Jan 1 – April 14	Spring May 1 – June 30	Summer July 1 – Sept 30
Legal	550	550	1200	1500
Sublegal	400	400	400	Project Ended
Overall total	Legal Sublegal	3800 1200		

Sport Fish Sampling Design

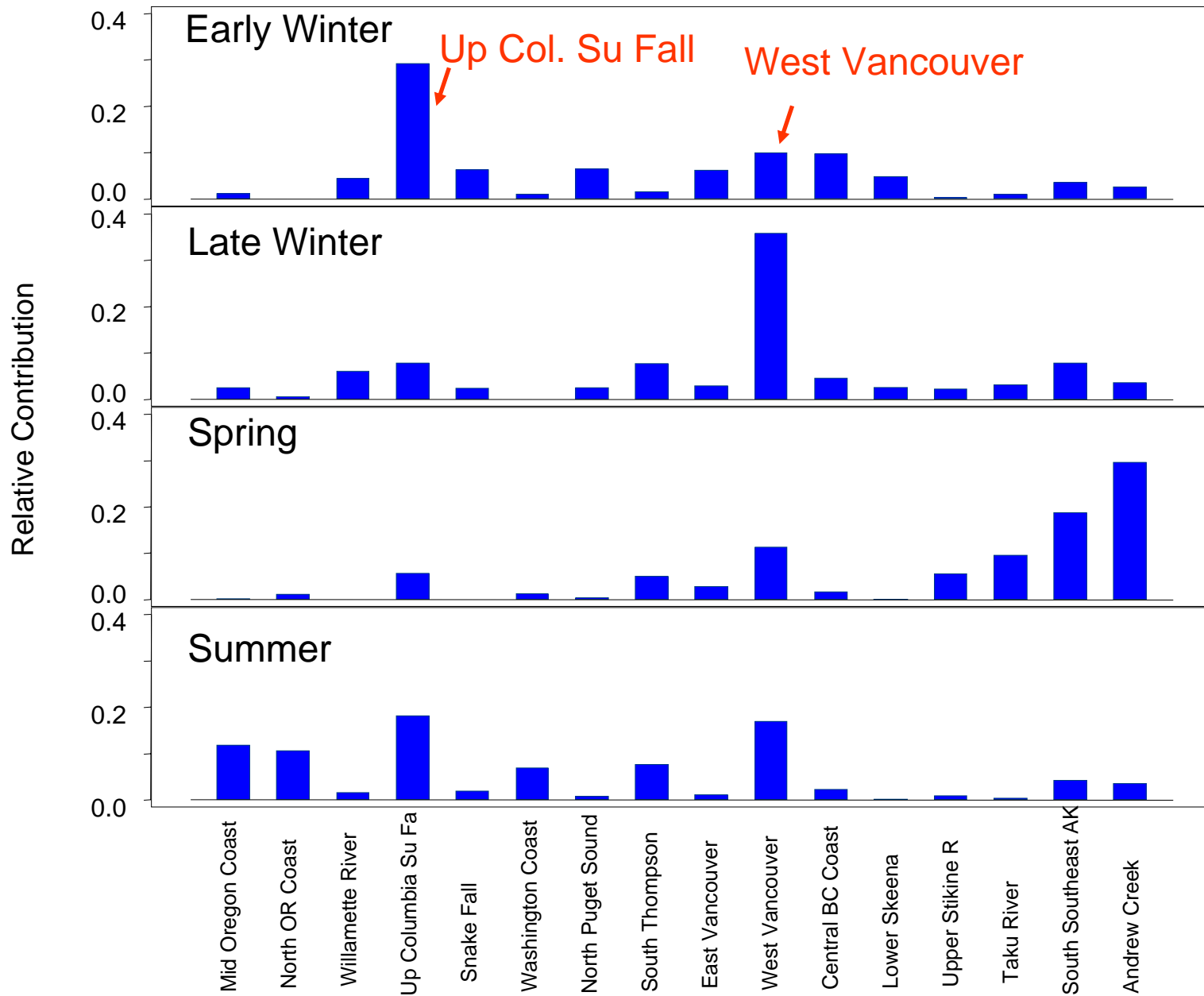
11 Ports

Size	Northern Inside	Northern Outside	Central	South
Legal	400	500	300	300
Sublegal	110	200	80	210
Overall totals	Legal Sublegal	1500 600		

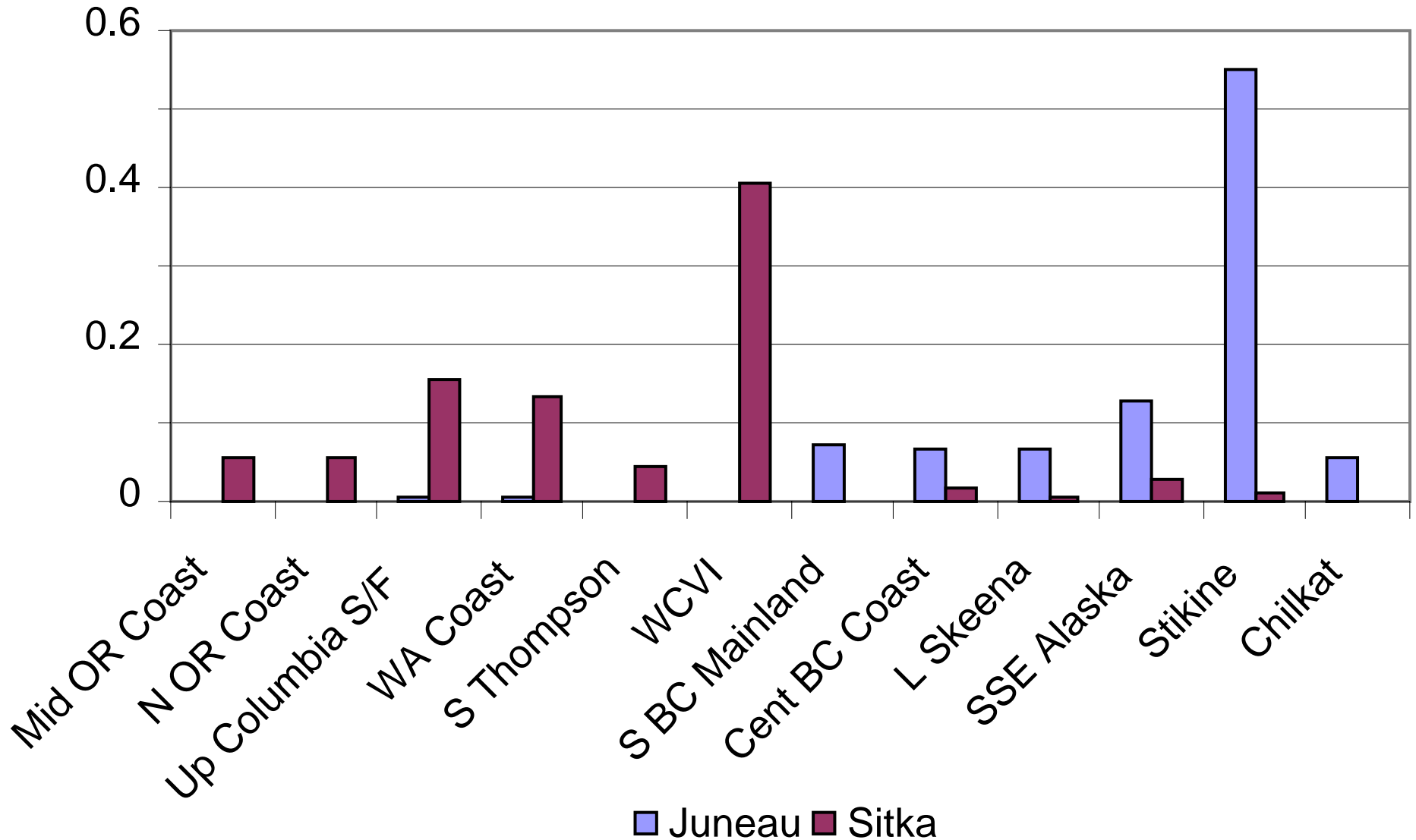
Gillnet Fishery Sampling Design

Area	District	Port	Goal	Weekly Sampling Rate
Stikine	108	Petersburg	440	40 fish per week
		Wrangell	<u>880</u>	80 fish per week
			1320	
Taku Inlet	111	Juneau	880	80 fish per week
Total			2200	

SE Alaska Troll Fishery 2004

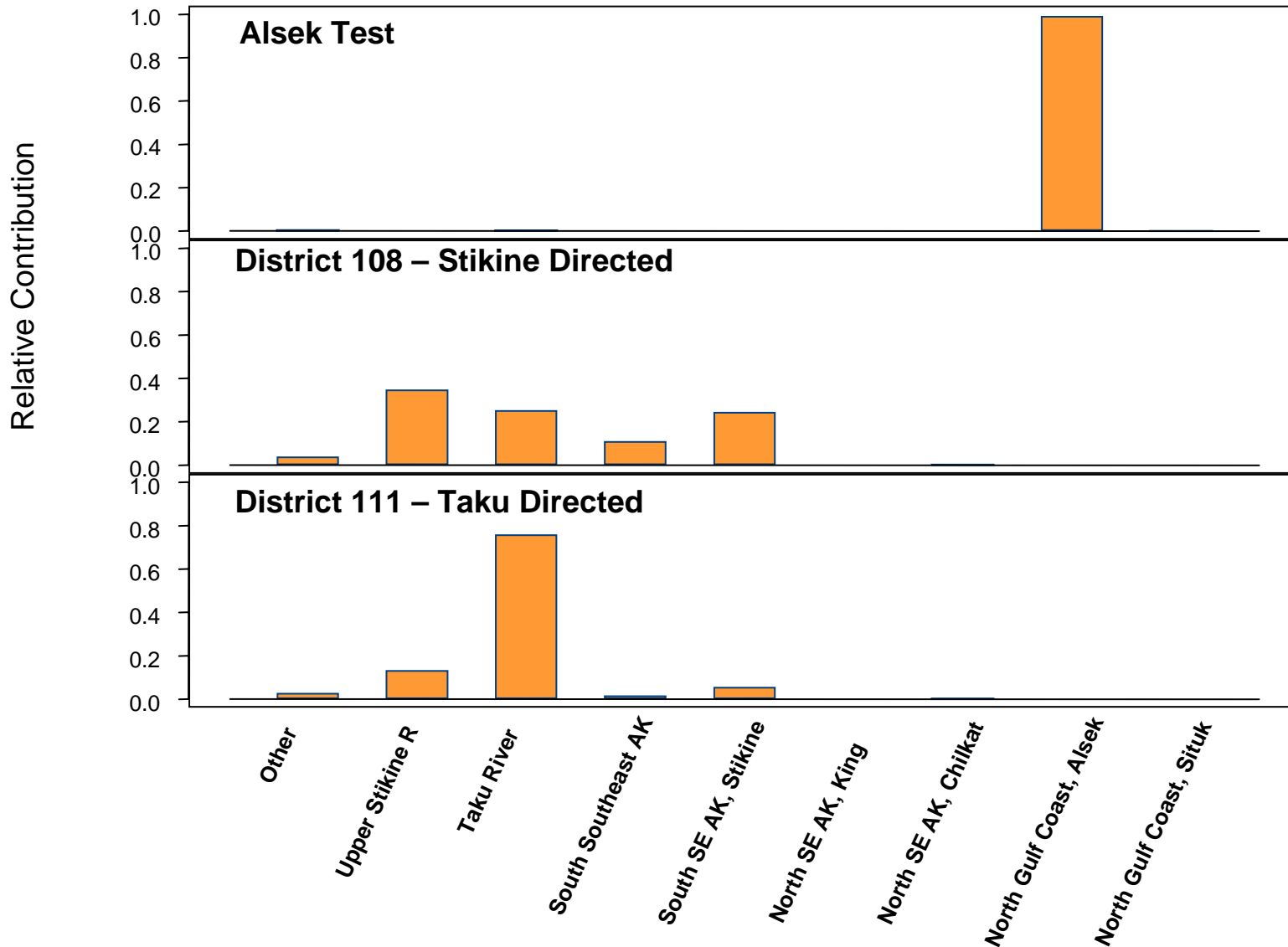


SE Alaska Sport Fishery 2004 after July 1



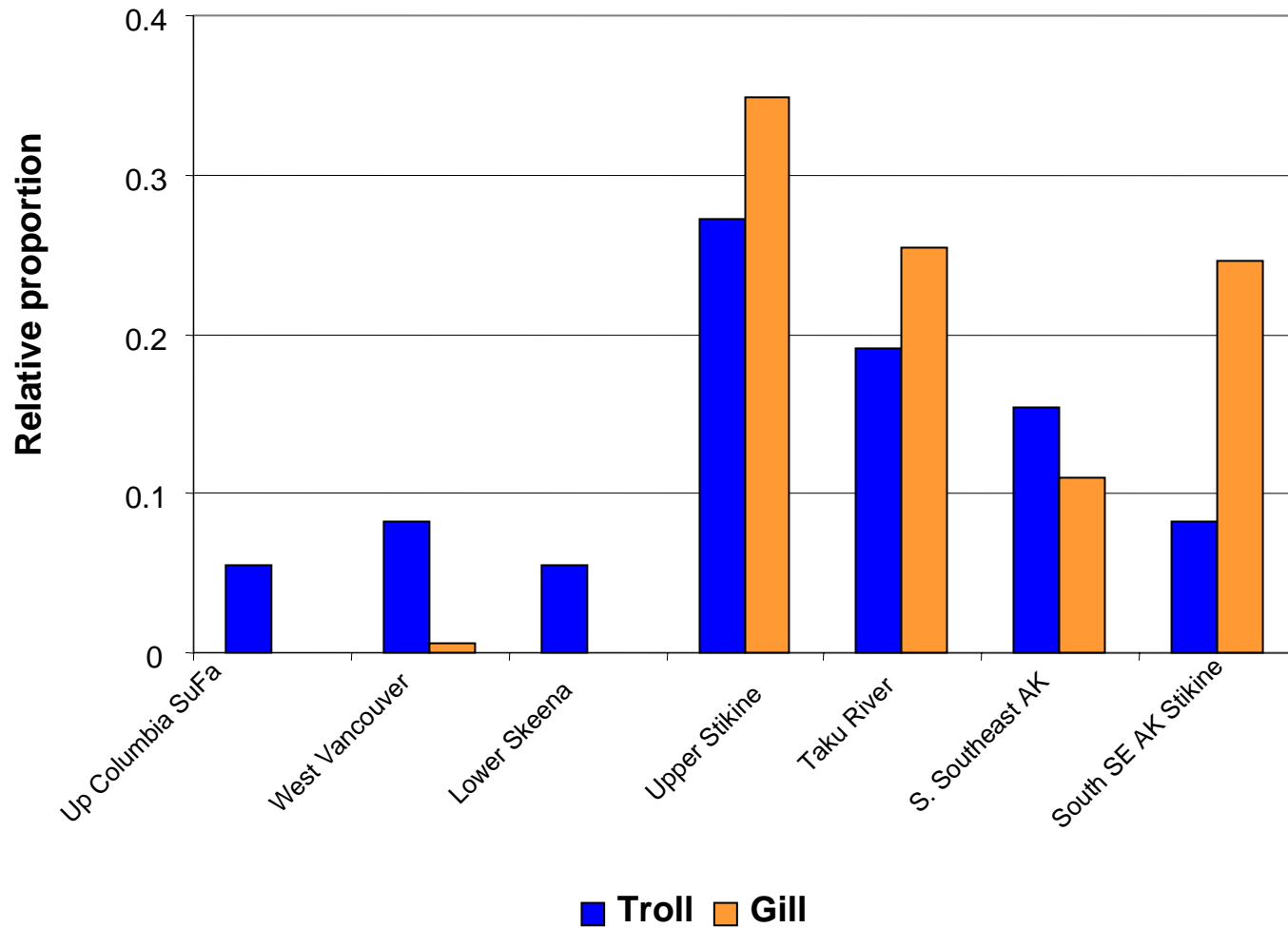
Local Gillnet Fisheries

2005



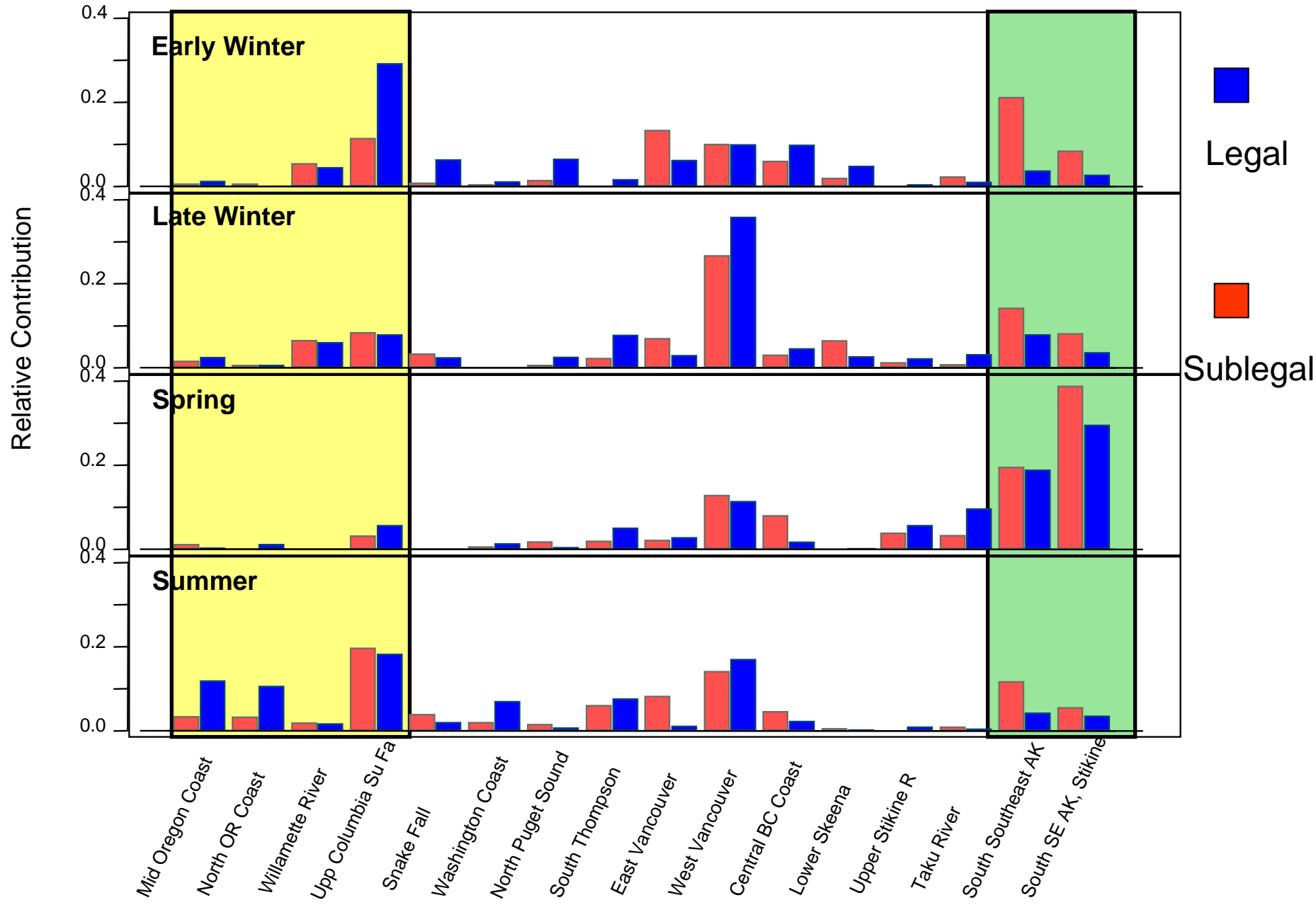
District 108 Directed Fishery - Stikine

2005 Troll vs. Gillnet



Legal and Sublegal Composition - Troll Fishery

Within 2004 Across Fisheries



ADFG Priorities

- Database development
 - Web accessible PSC GAPs database
 - Within laboratory—LOKI
- Development of high resolution genotyping
 - With Kalinowski/Taper from Montana State Univ.
 - *“Statistical methods to improve genetic stock identification” funded by US LOA*
- Conversion to SNPs
 - ADFG has converted all GSI applications to SNPs except SEAK Chinook
 - Pacific Rim Chinook, chum, sockeye
 - Improve automation, multiplexing, reduced costs

LOKI—Within Laboratory Database

- Control and error checking for:
 - Loci
 - Alleles
 - Collections/individuals
- Directly imports from multiple instruments
 - NO “cut and paste” or manual alignments
- Built in QC abilities
 - ADFG reruns 8% of scores
- Tracks metadata, tissue collections, extractions
- 5.5 million genotypes in ADFG database

File Security Importers Collection Utils Help

All Collections Editor

File Build Fish Database Help

CollectionPanel Fish and Tests Samplers

Speciespink salmon

Collection Date07-01-1991

Silly Code4P91ECN12180

RegionSouthcentral Alaska

LocationConstantine

Number of Samples100

QuadrantPWS - East

Latitude60.3609

Date Received01-01-1991

Longitude-146.6149

Collection ID4P91ECN1

☒ Verified Y/N

VerifiedByJMB

VerifiedDate01-25-2007

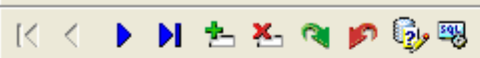
Comments

4 Replicate Samples taken, "1" on end of collectid added to make id unique
(Retro GPS, river mouth, not necessarily exact sampling location, 12/04, DM)

row 1Modified:false

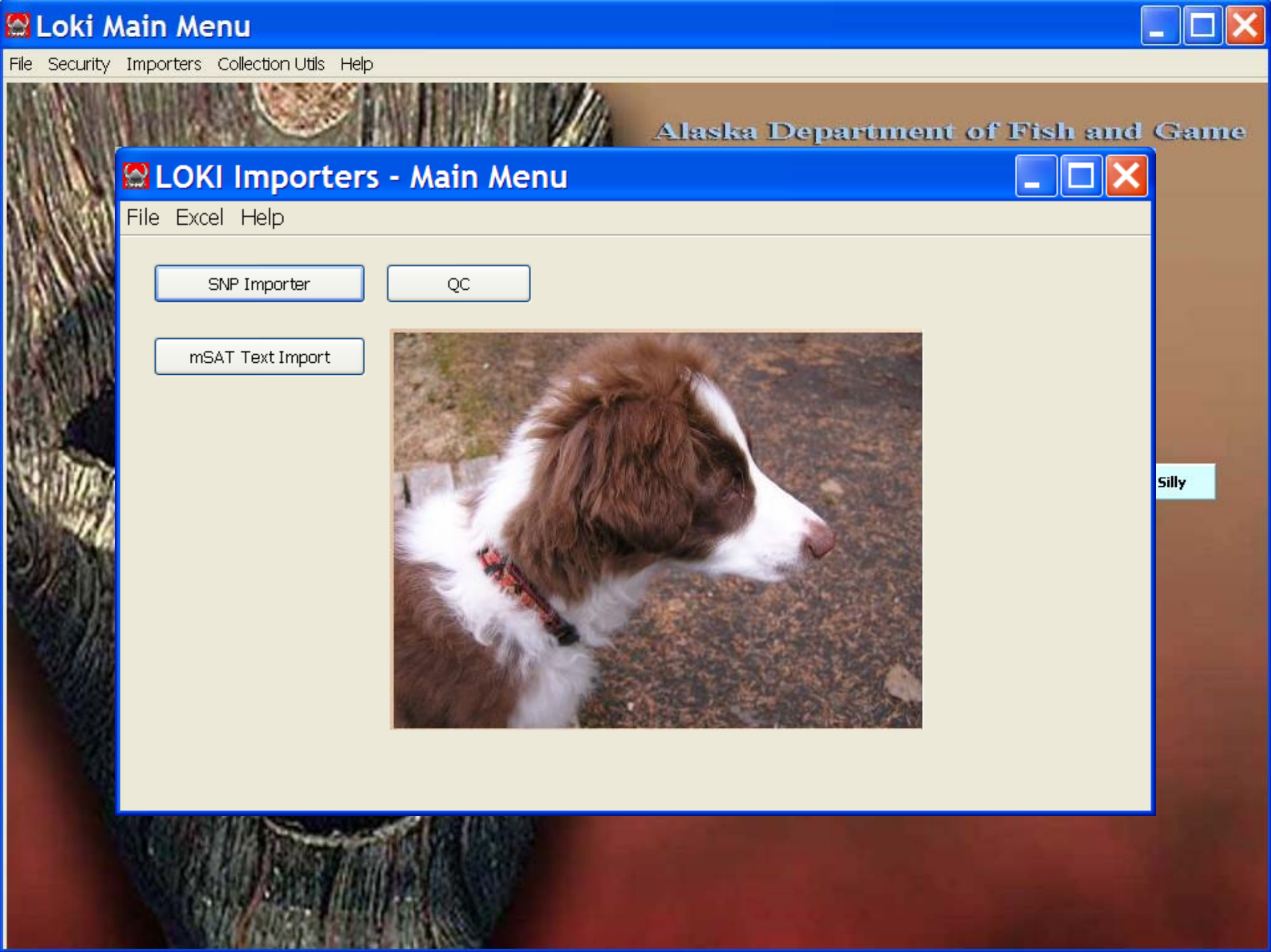
Locus and Allele Editor

File Database Help



Marker Alleles

Marker Name	<input type="text"/>	Citation	<input type="text"/>
Ploidy	<input type="text"/>		
Both Genotype 1	<input type="text"/>	Comment	<input type="text"/>
Both Genotype 2	<input type="text"/>		
Species	<input type="text"/>	Published Name	<input type="text"/>
Machine	<input type="text"/>	Concept Citation	<input type="text"/>
Dye	<input type="text"/>		
Lad...	<input type="text"/>	Validation Citation	<input type="text"/>
Genebank No.	<input type="text"/>	DB SNP No.	<input type="text"/>



Loki Main Menu

File Security Importers Collection Utils Help

Alaska Department of Fish and Game

LOKI Importers - Main Menu

File Excel Help

SNP Importer

QC

mSAT Text Import



Silly

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- Conversion to SNPs nearly complete
 - GSI applications converted to SNPs except SEAK Chinook
 - Highly automated, cost-efficient
 - Low error rate

ADFG Priorities

Development of high resolution genotyping

- PIs Kalinowski/Taper from Montana State Univ., Templin ADFG
 - *“Statistical methods to improve genetic stock identification” funded by US LOA*
- Objectives
 - Refine methods to estimate contribution of low freq stocks
 - Identification of the source of estimation error in mixtures
 - Incorporate improvements into GMA computer program

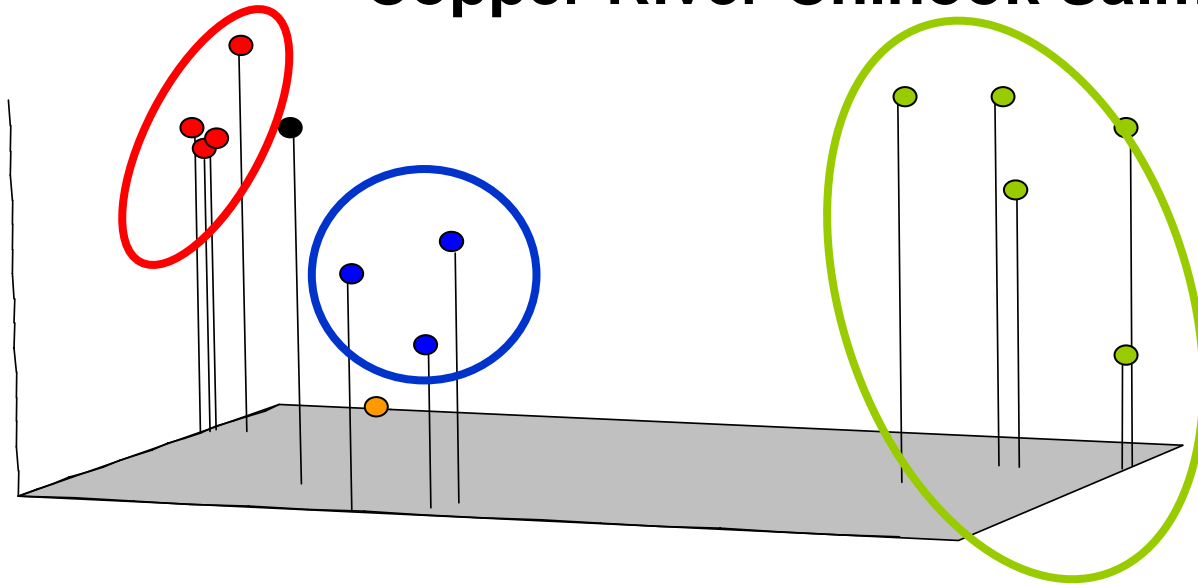
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 - Ease of standardization

SNPs

45 loci

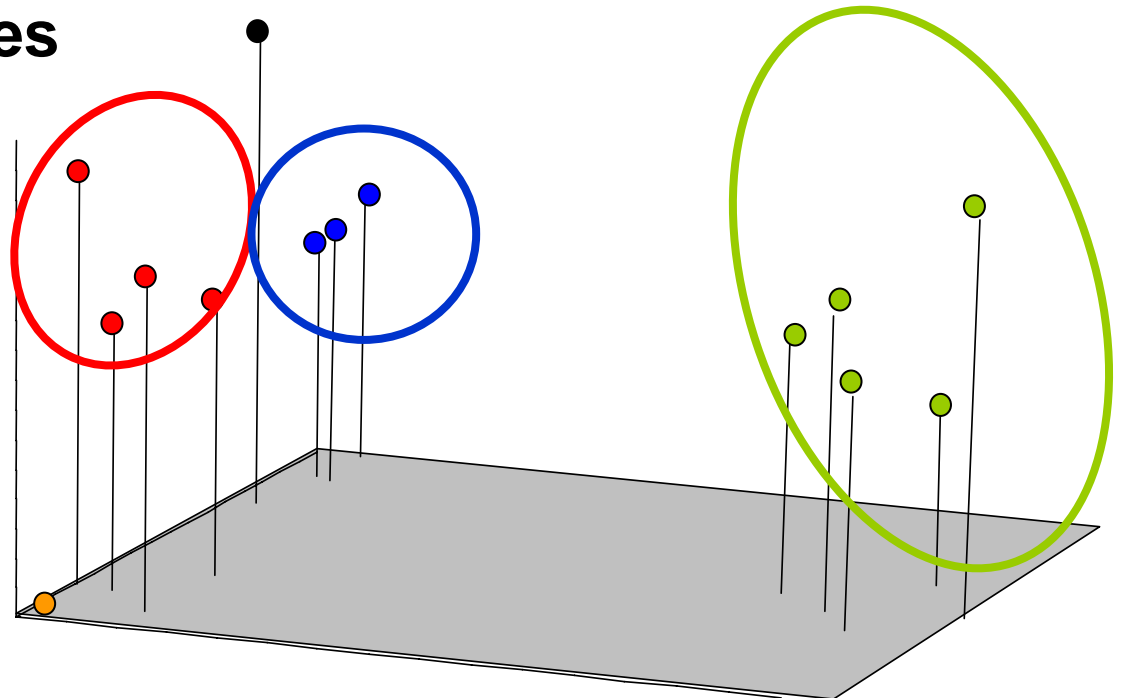
Copper River Chinook Salmon



Microsatellites

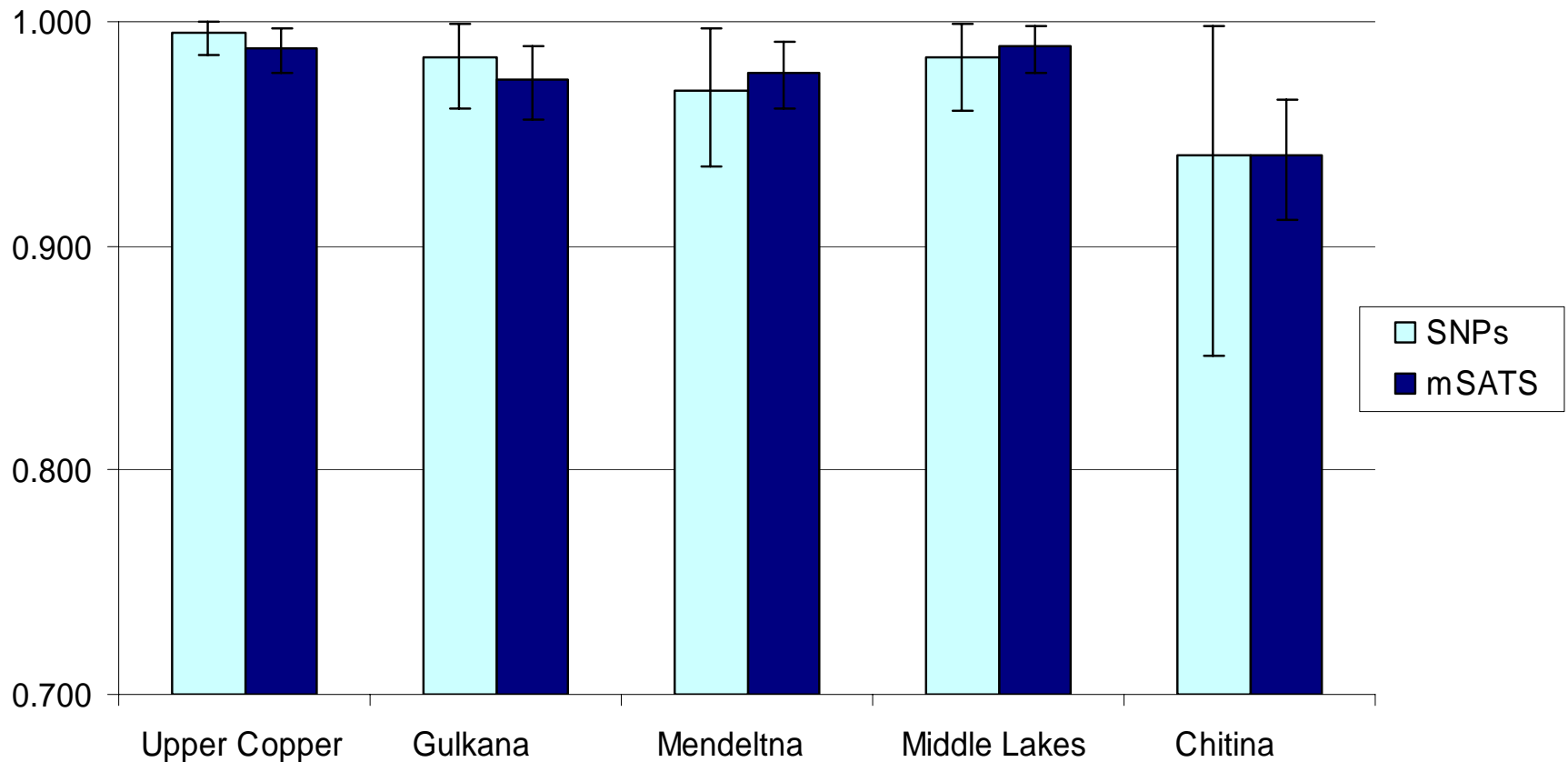
GAPs 13

High
concordance in
population
structure



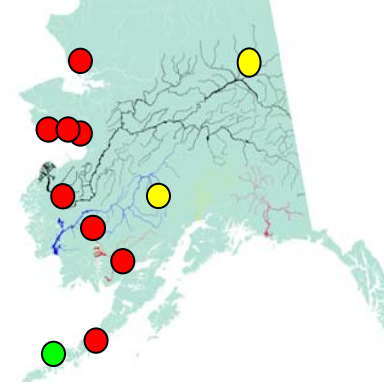
Genetic Markers for Stock Identification of Copper Chinook

100% Simulations By Region

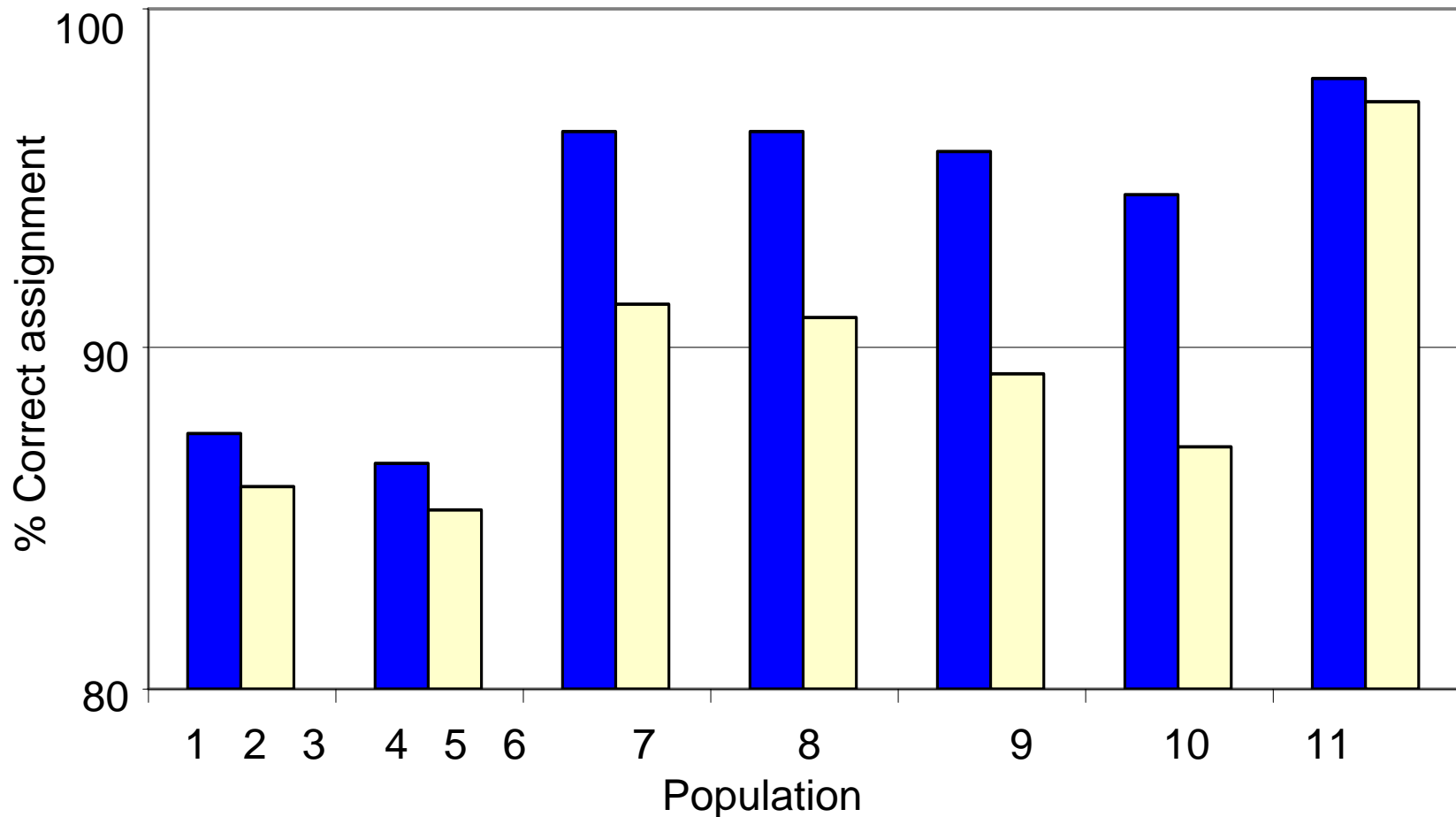


Alleles: 255 mSATS,
45 SNP loci

Accuracy of proportional assignments Chum Salmon



SNPs 48 alleles
Microsatellites 349 alleles



Cost of SNP Analyses

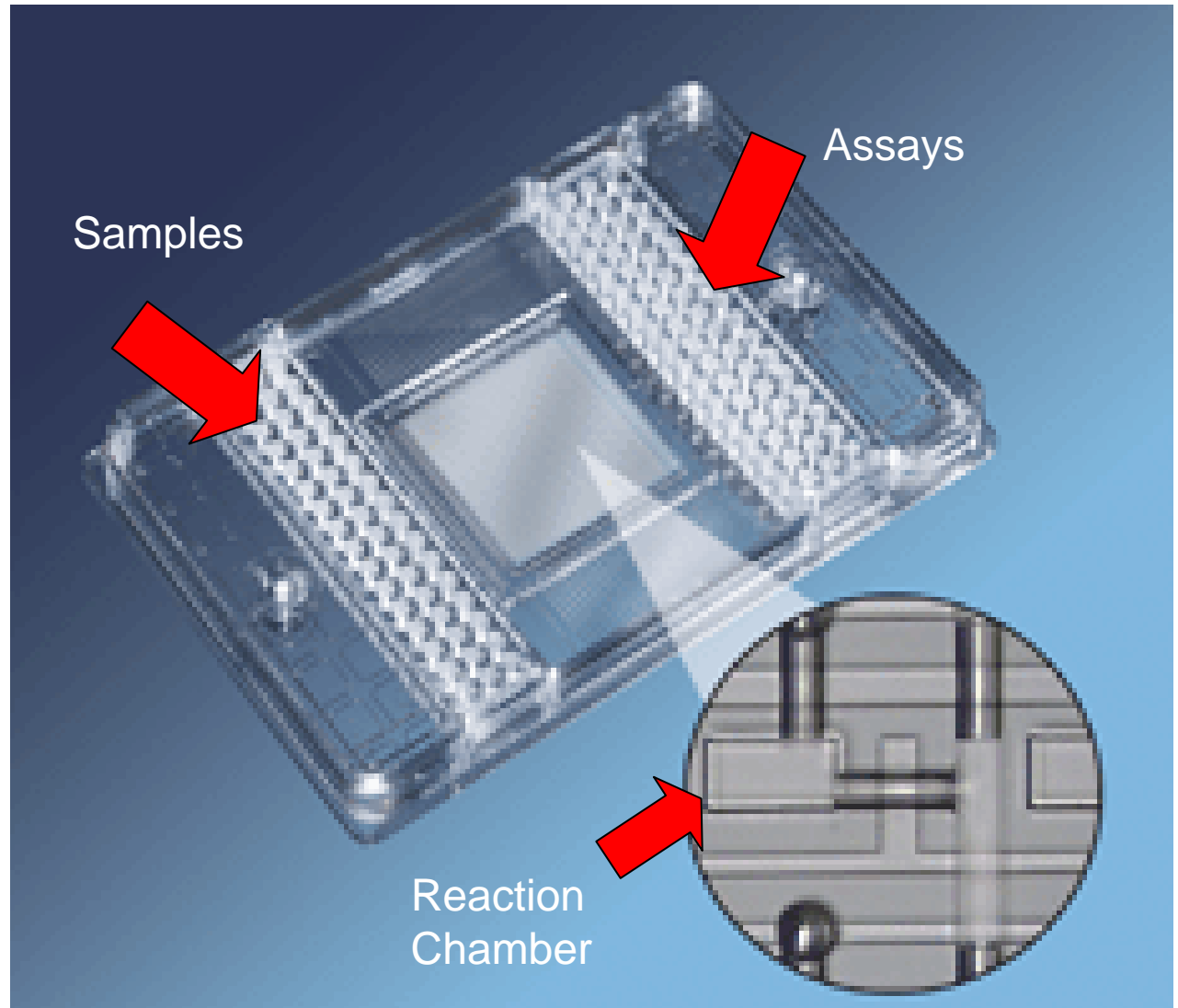
Reagents & Consumables

	Reaction volume	Cost/Genotype	Total cost (approx.)
96-well format, low volume purchasing	20 μ L	\$0.003 Plastics \$0.710 TaqMan \$0.827 Assay	\$1.540
96-well format, high volume purchasing	20 μ L	\$0.003 Plastics \$0.453 TaqMan \$0.220 Assay	\$0.676
384-well format, high volume purchasing	5 μ L	\$0.009 Plastics \$0.113 TaqMan \$0.055 Assay	\$0.177

48.48 Dynamic Array

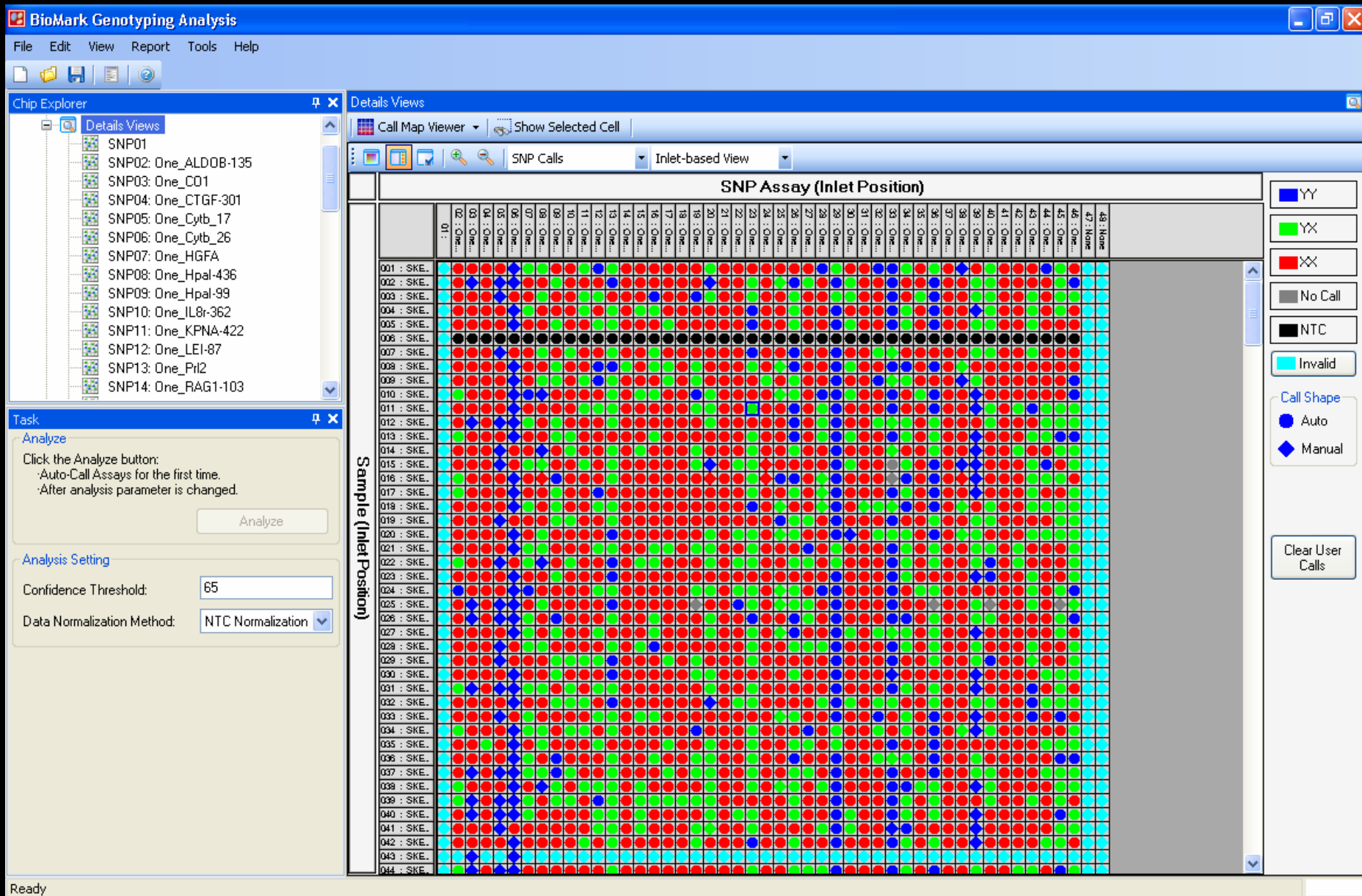
(48 samples X
48 assays)

10 nL reaction
volume



2304 wells

Biomark Chip by Fluidigm



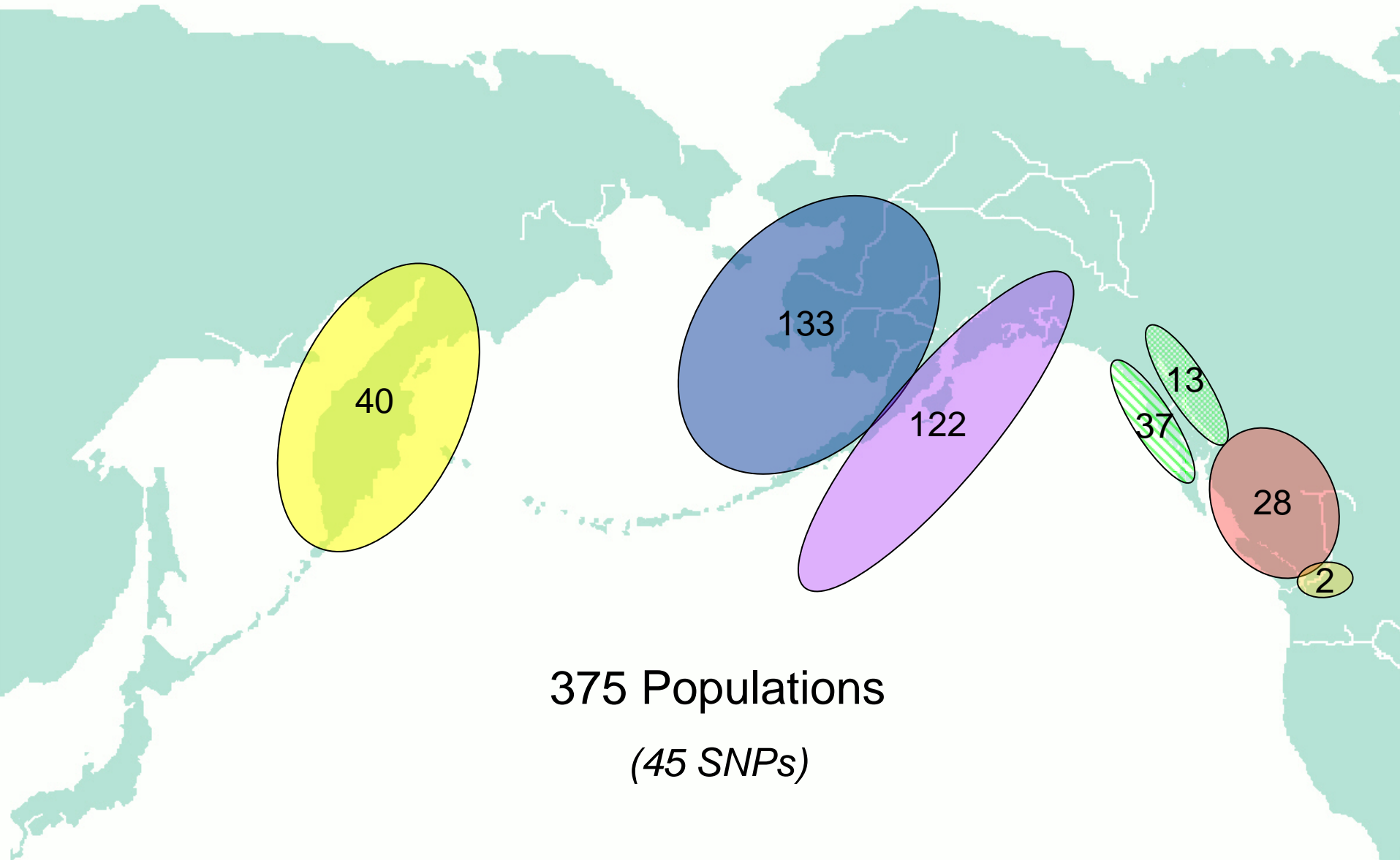
Biomark; 2304 wells

Cost of SNP Analyses

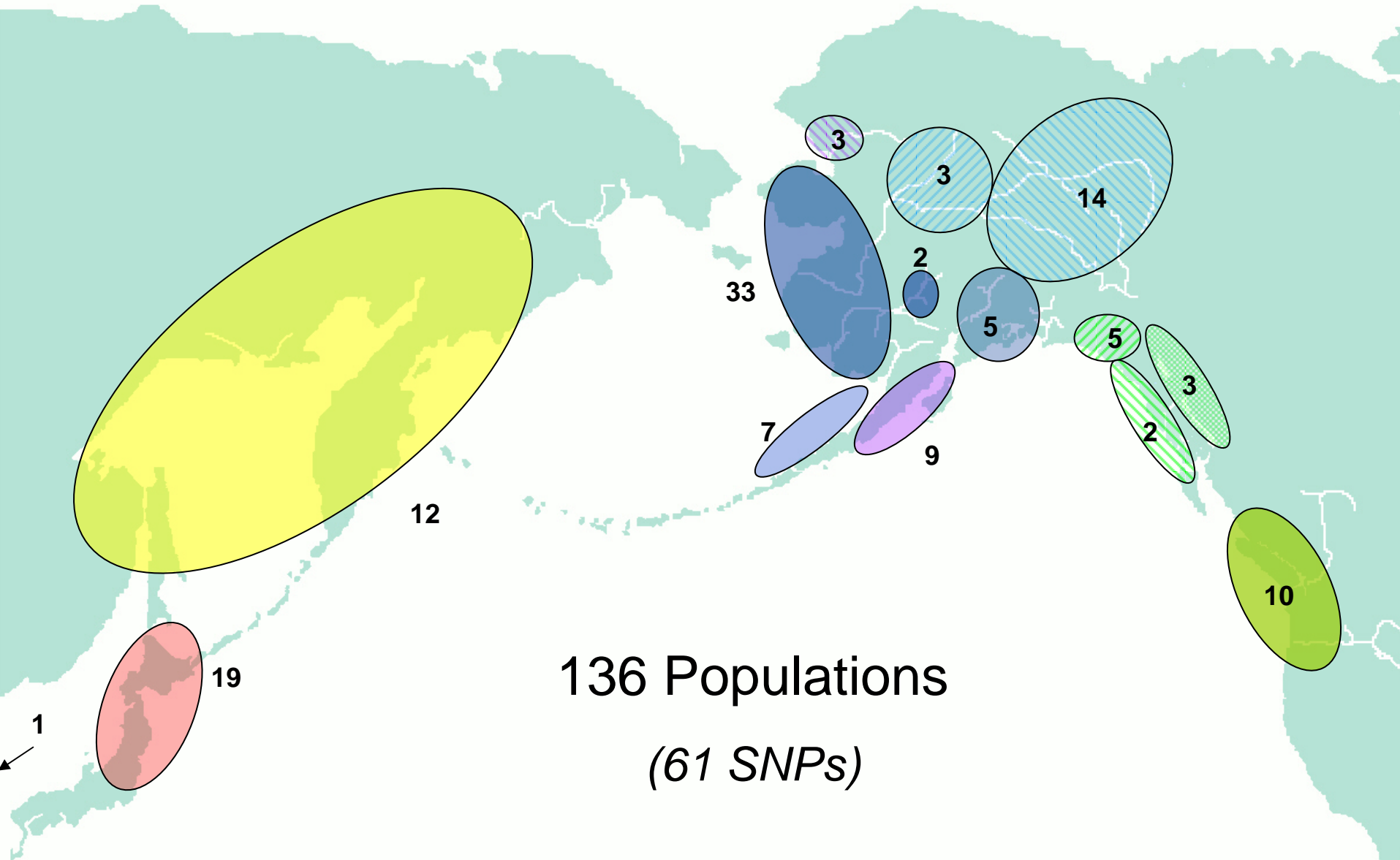
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Multiplex-48.48	10 nL	\$0.102 Chip \$0.002 TaqMan \$0.001 Assay	\$0.106

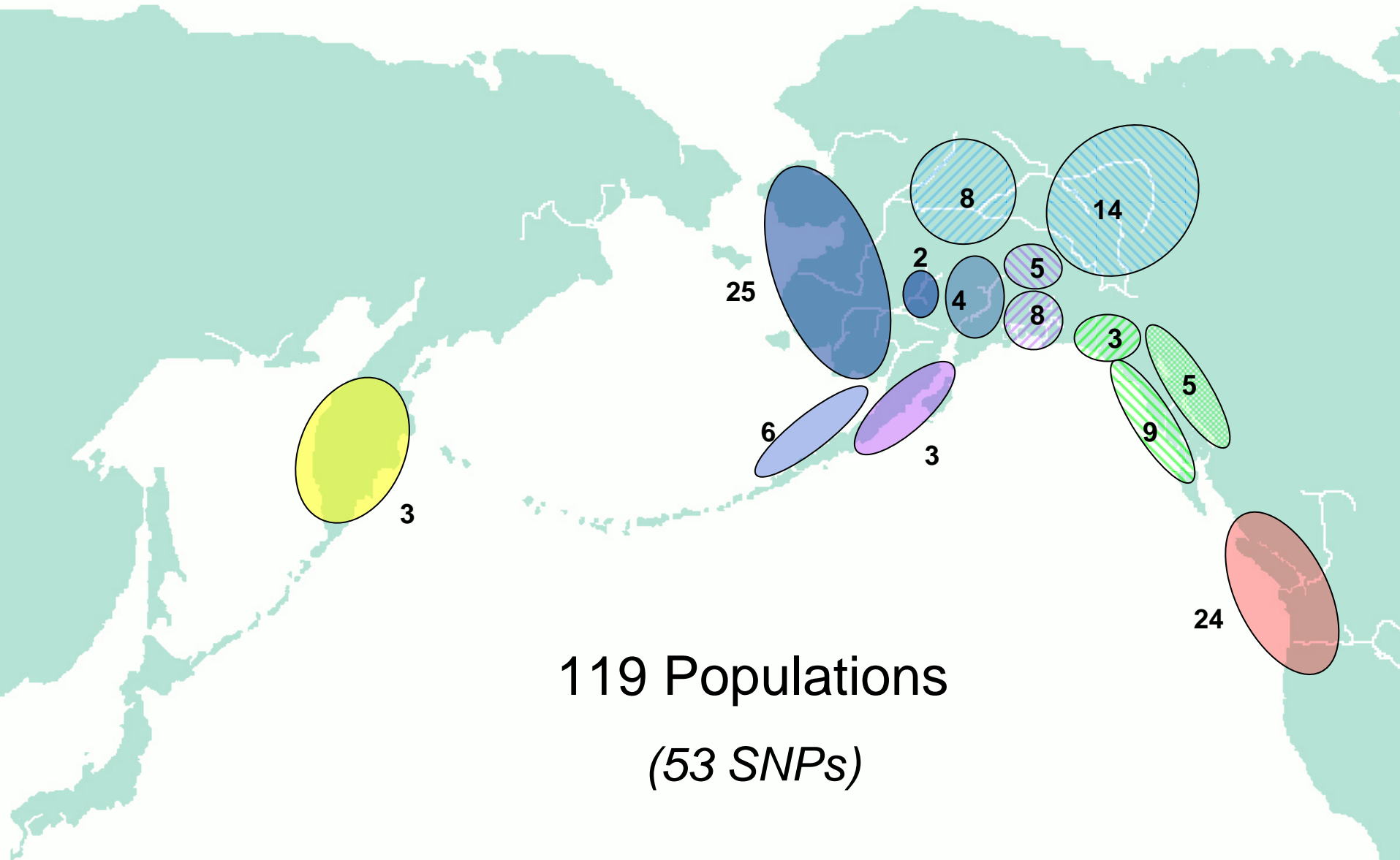
2007 Sockeye SNP Baseline



2007 Chum SNP Baseline

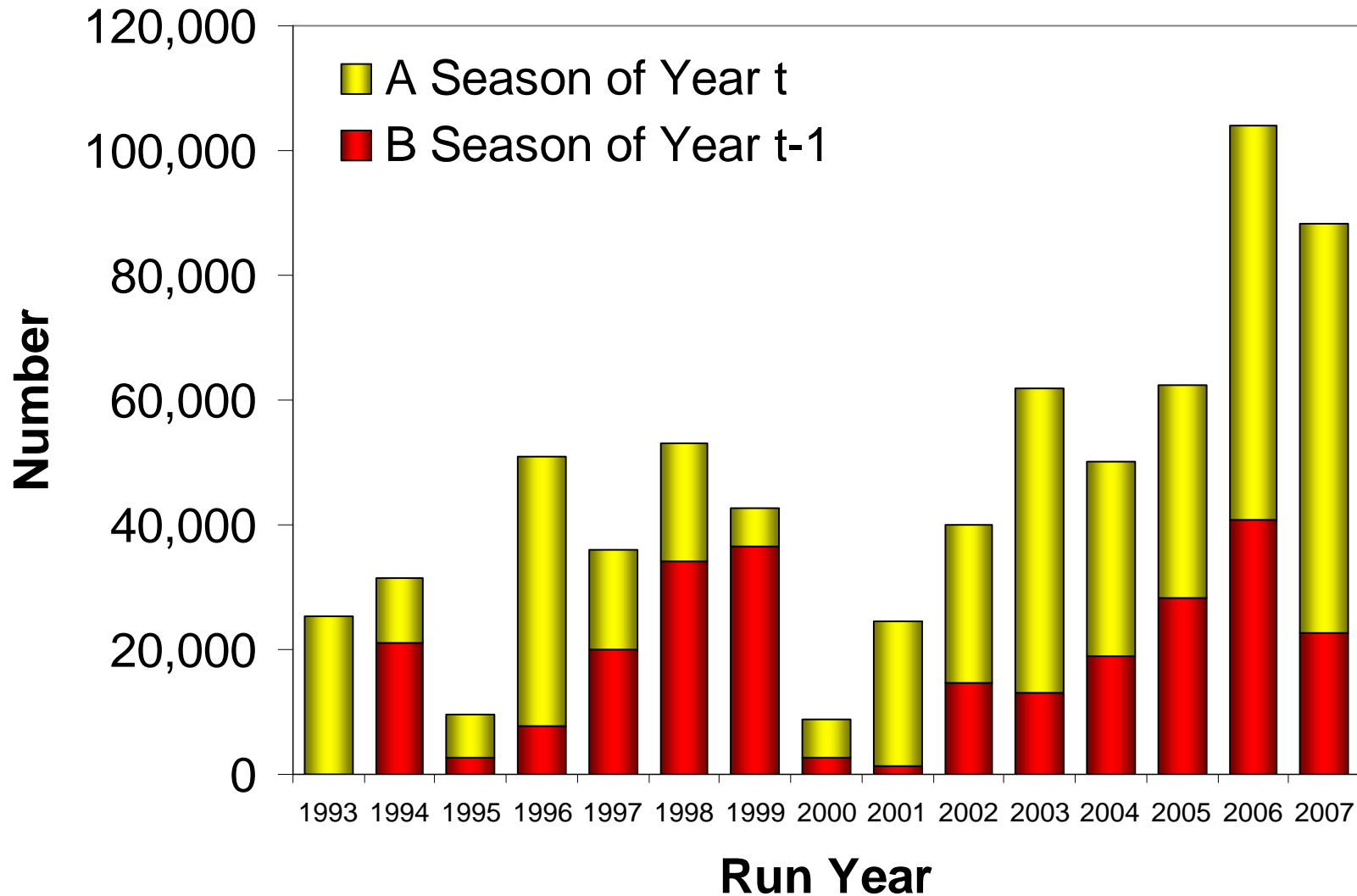


2007 Chinook SNP Baseline

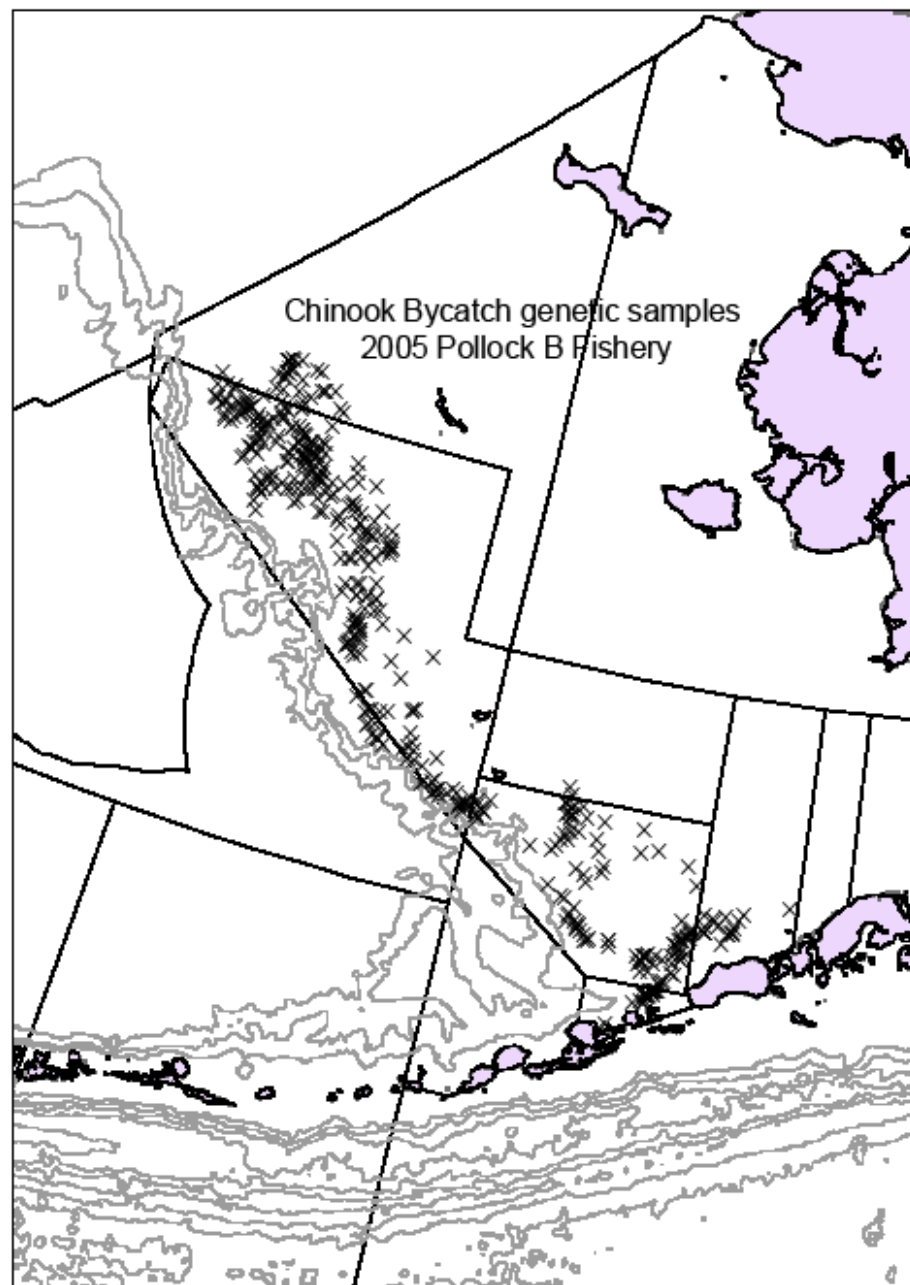


BSAI Bycatch of Chinook salmon

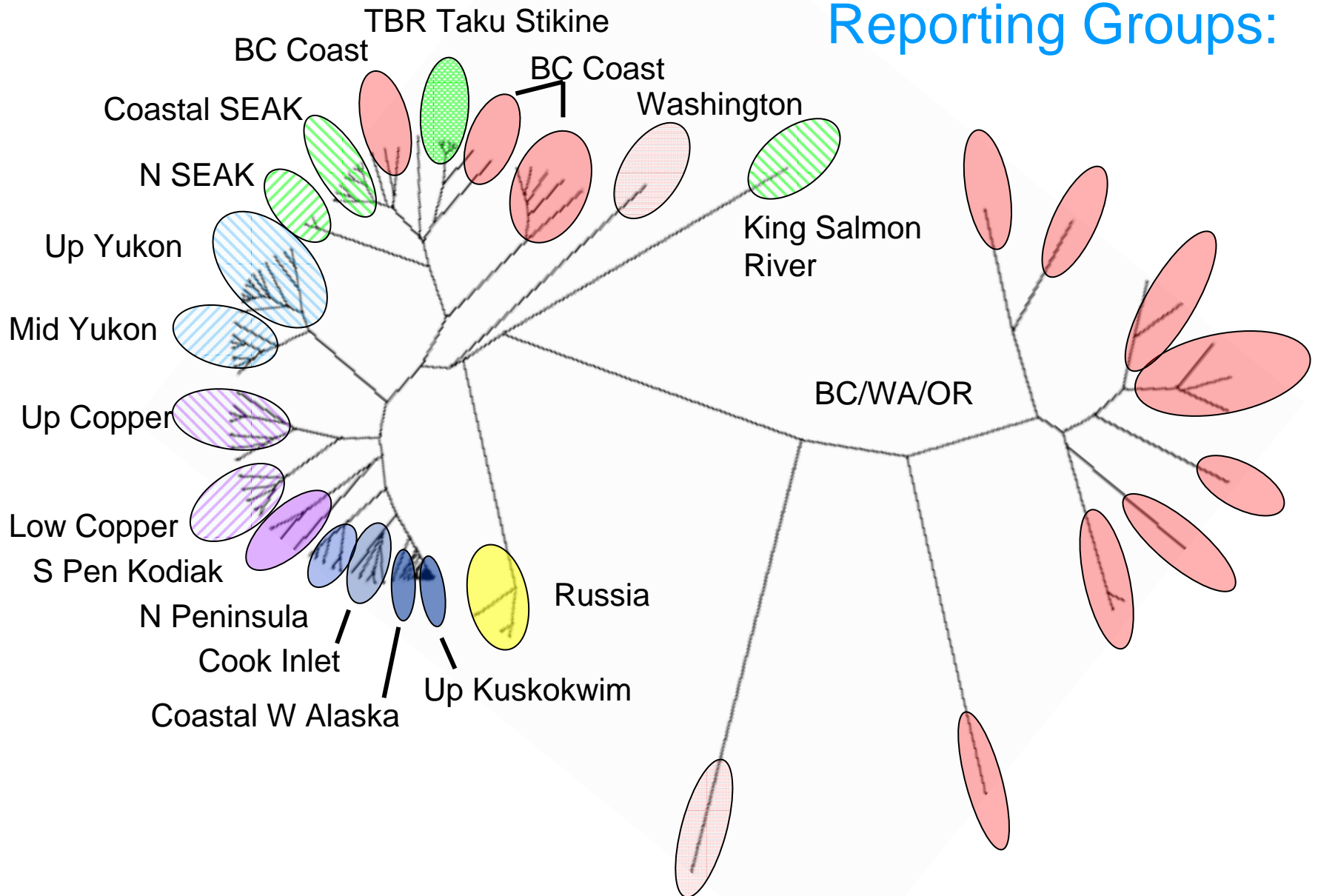
(by run year)



Origin of samples: 2005 Pollock B fishery

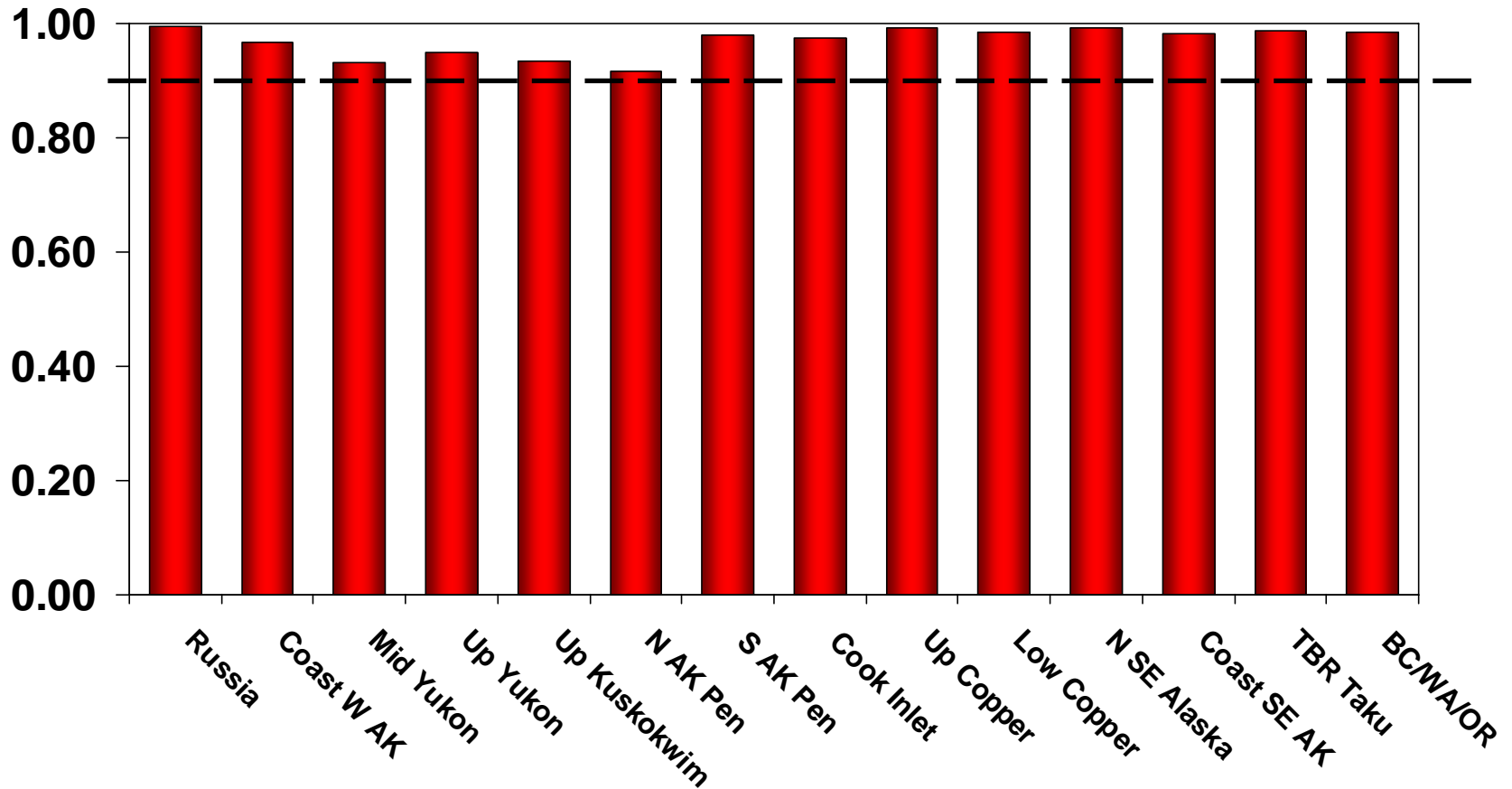


Broad-scale Reporting Groups:



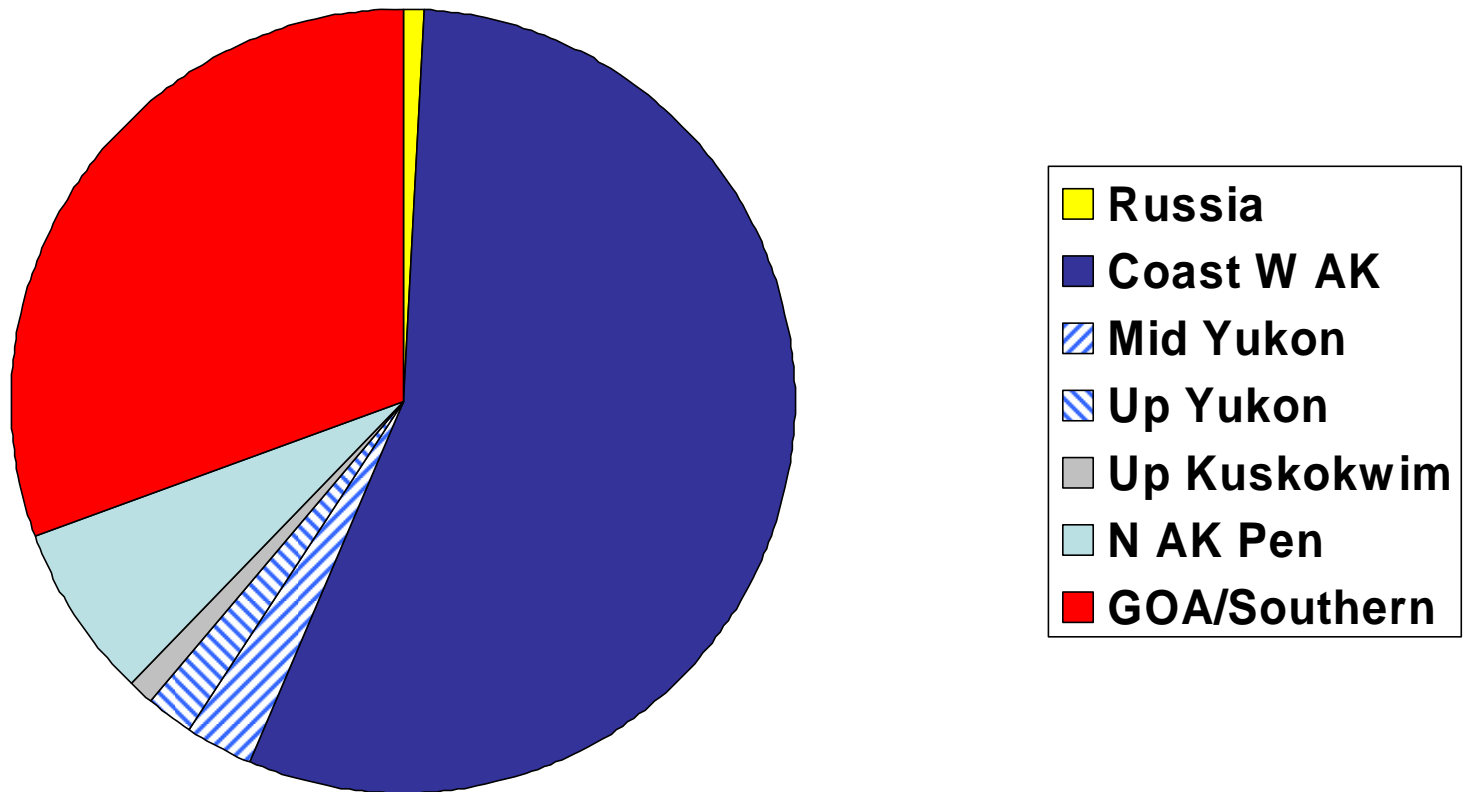
100% Simulations

(48 SNPs)



2005 B Season Estimates

(Composition of sample)



N = 887

GSI Applications in PSC ADFG Perspective

- Fishery sampling--extensive
 - Multiple strata---time, area, age, gear
 - Sample size adequate to provide resolution required by managers (control sampling error)
- Laboratory processing--highly automated, cost-efficient
 - Continued baseline improvement (control genetic error)
 - Very low genotyping error rate, extensive QC procedures
 - Robotics, barcoding, fast turnaround when needed
 - Coastwide labs will vary in processing power
 - Low throughput -- 10,000+ samples/year
 - High throughput (most efficient) -- 50,000+ samples/year
- Data processing
 - Statistical development to reduce effects of errors
 - Within- and among-laboratory databases

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GSI Applications in PSC

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Acknowledgements

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