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Deploying protected species tools via cloud computing

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November 18, 2015

Web-based Protected Species Toolbox and Cloud-based Platform for Running Tools

- **PIs:** Eli Holmes and Eric Ward (NWFSC)
- **Goal:** To develop both a website/content manager for protected species tools and browser-based tool interfaces.
- **Importance/Application:**
 - NMFS scientists develop many applications/tools to run sophisticated analyses in support of NMFS Regional Offices.
 - There is currently no cross-center platform to host these tools. Tools existence often unknown across groups.
 - Browser-based interfaces allow NMFS scientists to develop tools that can be run by others with only a browser.

Webpage



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Marine Mammals & Turtles

Ecosystems

Stock Assessments

Economics

Human Dimensions

Apps & Tech

Climate

Welcome to the Protected Species Toolbox

The National Protected Species Toolbox (NPST) provides access to a variety of modeling and statistical tools used to support the protection, conservation and recovery of **marine mammals** and **endangered/threatened marine life** under the responsibility of **NOAA's** National Marine Fisheries Service (NOAA Fisheries Service, or **NMFS**), under the **U.S. Department of Commerce**. These tools are developed and maintained by individual NMFS science centers.

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Project pages

ROSE — [Download files](#)

Project members: e2holmes eric.ward howard.coleman

Center	System	Category
NWFSC	Marine Mammals	Demographic Analysis

Version:	Language:	Keywords:
1.0	R, jags	risk assessment

Summary

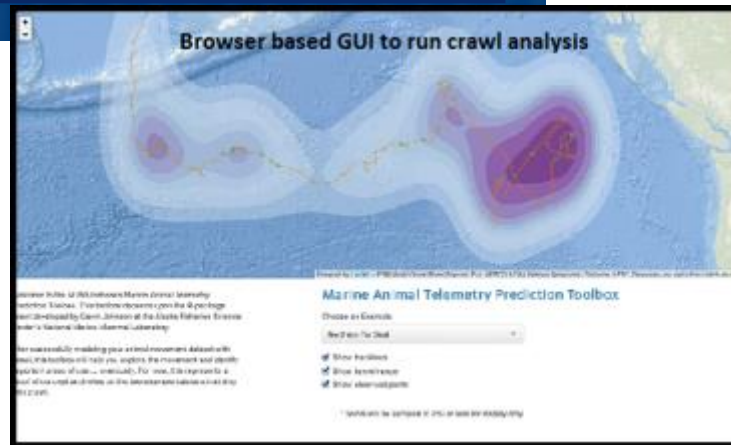
Several of the smaller populations that NMFS monitors have very detailed demographic data, where individual birth and deaths are known, in addition to the reproduction of each female in each year. The Resident Orca Salmon Emulator (ROSE) model is a tool estimate how survival and birth rates of these small populations changes over time, and whether any of these rates is affected by external drivers (climate, prey, etc). Small adjustments to these drivers may be important if the external driver is a prey species that is also commercially fished. Most recently, this tool has been applied to estimate how altering fishing levels of Chinook salmon may impact the viability and growth of endangered Southern Resident killer whales.

How is ROSE used in analyses related to protected species management?

ROSE is used to understand how the viability of Southern Resident Killer Whales is related to covariates such as prey. ROSE was used in the bi-lateral workshops investigating the relationships between Chinook salmon fishing and Southern Resident Killer Whale viability (report) and was used in the 2013 NOAA Technical Memorandum Estimating the Impacts of Chinook Salmon Abundance and Prey Removal by Ocean Fishing on Southern Resident Killer Whale Population Dynamics.

[View ROSE details](#)

Browser-based GUIs



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e2holmes

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Forecasting and Simulation	Center	System	Lang	Vers
Count-based PVA	NWFSC		matlab	1.0
DARTER	NWFSC		Excel	
Selective Harvest Calculator (SHC)	NWFSC	Anadromous Fish	Java	1.0.5
Species Life-cycle Analysis Modules (SLAM)	NWFSC	Anadromous Fish	Java	
Viability and Risk Assessment Procedure (VRAP)	NWFSC	Anadromous Fish	R	1.0
Demographic Analysis				
ROSE	NWFSC	Marine Mammals	R, jags	1.0
Salmon Population AnalyZer (SPAZ)	NWFSC	Anadromous Fish	Java	1.3.4
Analysis of Movement Data				
Genetic Analysis				
gsi_sim	SWFSC	Anadromous Fish		
Diet and Stable Isotope Analysis				
Statistical Packages				
MARSS	NWFSC		R	3.13

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Forecasting and Simulation

Count-based PVA
DARTER
Selective Harvest Calculator (SHC)
Species Life-cycle Analysis Modules (SLAM)
Viability and Risk Assessment Procedure (VRAP)

Center **System**

NWFSC
NWFSC
NWFSC Anadromous Fish
NWFSC Anadromous Fish
NWFSC Anadromous Fish

Demographic Analysis

ROSE
Salmon Population AnalyZer (SPAZ)

NWFSC Marine Mammals
NWFSC Anadromous Fish

Analysis of Movement Data

Genetic Analysis

gsi_sim

SWFSC Anadromous Fish

Diet and Stable Isotope Analysis

ROSE

PST Admin: [Remove 'ROSE' from P](#)

Project information

Published : [Remove](#)

Project members
e2holmes howard.coleman eric.ward
[Manage users](#)

Sharing: Public Center: NWFSC System: Marine Mammals Category: Demographic Analysis

Summary

Several of the smaller populations that NMFS monitors have very detailed demographic data, where individual birth and deaths are known, in addition to the reproduction of each female in each year. The Resident Orca Salmon Emulator (ROSE) model is a tool estimate how survival and birth rates of these small populations changes over time, and whether any of these rates is affected by external drivers (climate, prey, etc). Small adjustments to these drivers may be important if the external driver is a prey species that is also commercially fished. Most recently, this tool has been applied to estimate how altering fishing levels of Chinook salmon may impact the viability and growth of endangered Southern Resident killer whales.

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ROSE is used to understand how the viability of Southern Resident Killer Whales is related to covariates such as prey. ROSE was used in the bi-lateral workshops investigating the relationships between Chinook salmon fishing and Southern Resident Killer Whale viability ([report](#)) and was used in the 2013 NOAA Technical Memorandum [Estimating the Impacts of Chinook Salmon Abundance and Prey Removal by Ocean Fishing on Southern Resident Killer Whale Population Dynamics](#).

Detailed description

To install: Download the zip file for the package ([click to download](#)). Then use the devtools package in R.

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7 project downloads
26 file downloads
[View downloads](#)



NWFSC Data Explorer : Conservation Biology Program

These R Shiny applications are based on packages developed in conjunction with the Conservation Biology Program of the Northwest Fisheries Science Center.

To run an application, or to learn more about it, click its link.

To return to this page, click "Display apps" in the bottom right-hand corner of the application display.

Shiny applications

[agTrend](#) : Estimating trends of aggregated abundance

[ROSE](#) : Resident Orca Salmon Emulator

[VRAP](#) : Viability Risk Assessment Procedure & Rebuilding Exploitation Rates

agTrend: AFSC Steller sea lion trend analysis and data visualization

<https://dataexplorer.northwestscience.fisheries.noaa.gov/nwc/agTrend/>



agTrend

Select data set | **Plot data** | Fit model | Abundances

Select regions and sites:

Regions:

- C ALEU
- E ALEU
- W ALEU
- C GULF**

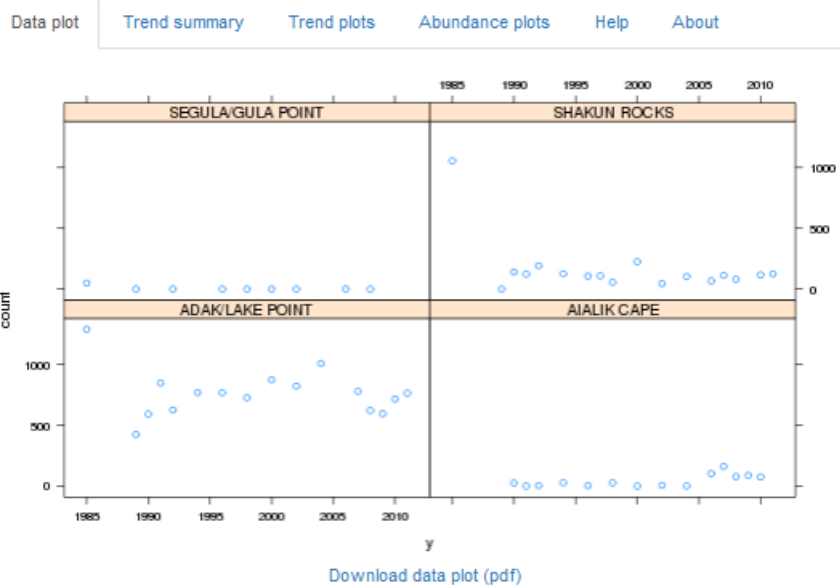
Available sites

- SEAL ROCKS
- SEAL ROCKS (KENAI)
- SEGUAM/SADDLERIDGE
- SEGUAM/TURF POINT
- SEGULA/CHUGUL POINT
- SEMISOPOCHNOV/PETREL
- SEMISOPOCHNOV/POCHNOI
- SEMISOPOCHNOV/SW KNOB
- SHAW
- SHEMYA
- SILAK

Selected sites


- ADAK/LAKE POINT
- AIALIK CAPE
- SEGULA/GULA POINT
- SHAKUN ROCKS**

Plot data



VRAP: NWFSC Tool for estimation of impact of harvest on listed salmonids

<https://dataexplorer.northwestscience.fisheries.noaa.gov/nwc/VRAP/>



VRAP 1.1

Data input Results Help About

Step 1: Input data

Choose a demo file (click to paste data instead)

Select demo file:
Bev-Holt, no covariates, ER

estimated time for this file and NRRuns = 0 seconds

Step 2: Choose (or change) number of runs (NRRuns) per simulation

- 1
- 10
- 100
- 1000
- Use .rav NRRuns

Step 3: Run VRAP. Button appears after step 1 completed.

Run VRAP with selected file and NRRuns

Download the example .rav files:
Bev-Holt, no covariates, ER

Download Example File

Table of target exploitation rate versus extinction risk

VRAP 1.1

Data input Results Help About

R&PVIABILITY (R) Version 1.0 Date:2015-11-16

Title: ER example of Beverton-Holt VRAP with no marine survival or stream flow variability

Input File: demofiles/exampleB2ER.rav

Copy of Input File:demofiles/exampleB2ER.rav

Basic Simulation Input Parameters:
of Years=25 # of Reps=1 HR Conv.Crit=0.001 Seed0
Range start 0 end 1.89189189189189 by 0.0675675675675676

Stock Recruit Function Input Parameters:
Function Type: BEV2
Recruits = 1/[(1/b) + 1/(a*Spawners)]
a=productivity=9.27599153996063 b=maxrecruits=3702.14272977426

Stock-Recruit Error Parameters (gamma distr.) [R=f(s)+e]:
A=0.967637142385205 B=1.09275572895811

Depensation at escap:400 QET:63fraction of depensation at QET0

Fishery Regime Parameters:
Base ER = 0.37

Management Variability Parameters:
Gamma A=65.2846 Gamma B=0.0158
mean=1.03223468 var=0.016925107944

AEQ for age class
Age2 AEQ =0.584052132848285
Age3 AEQ =0.825713254893611
Age4 AEQ =0.968504491293936
Age5 AEQ =1
recruits Ab Age 1= 0.175215639854485

Regime Evaluation Parameters: QET = 63
Lower Escapement Level (LEL)=416
Upper Escapement Level (UEL)=1040
Max Return (under average variability) =3702.14272977426

SUMMARY STATISTICS
All statistics are averaged over repetitions

. param.	. b	Total-Exploit.-Rate	Escapement				#fish	truns	tYrs	1st LastYrs	pop_size.
			TgtER	CYtER	BYtER	Mort. extinct					
3702	0.00	0.000	0.000	0	0.0	0.0	100.0	722	1565		
3702	0.03	0.027	0.026	72	0.0	0.0	100.0	707	5322		
3702	0.05	0.051	0.051	111	0.0	0.0	100.0	700	2384		
3702	0.08	0.081	0.079	200	0.0	0.0	100.0	683	2243		
3702	0.10	0.100	0.098	214	0.0	4.0	100.0	677	4294		
3702	0.13	0.126	0.124	446	0.0	0.0	100.0	658	1807		
3702	0.15	0.152	0.148	589	0.0	0.0	100.0	657	4447		
3702	0.18	0.189	0.185	596	0.0	0.0	100.0	624	2867		
3702	0.20	0.206	0.202	679	0.0	0.0	100.0	608	1400		
3702	0.23	0.222	0.216	918	0.0	0.0	100.0	604	4149		

PVAwidget: NWFSC Tool for estimation of extinction risk from count data

<https://dataexplorer.northwestscience.fisheries.noaa.gov/nwc/PVAwidget>

PVAwidget

Data PVA

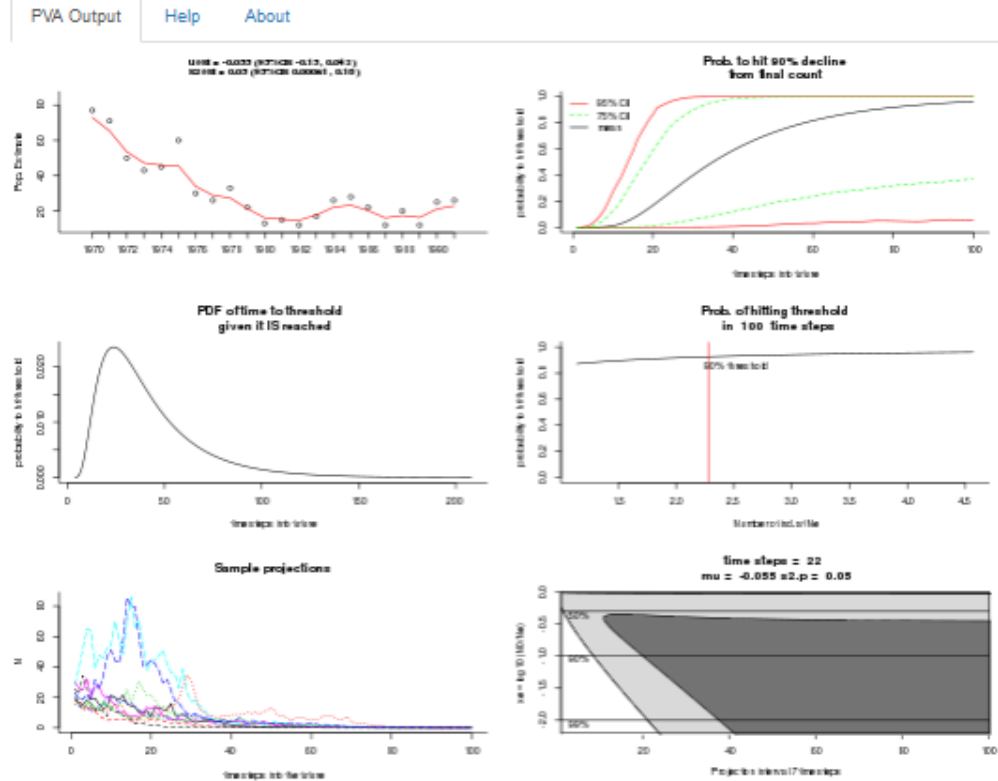
Paste CSV data

```

"","Year","Count"
"1",1970,77
"2",1971,71
"3",1972,60
"4",1973,43
"5",1974,45
"6",1975,60
"7",1976,30
"8",1977,26
"9",1978,33
"10",1979,22
"11",1980,13
"12",1981,15
"13",1982,12
"14",1983,17
"15",1984,26
"16",1985,28
"17",1986,22
"18",1987,12
"19",1988,20
"20",1989,12
"21",1990,25
"22",1991,26
    
```

Load CSV data

Data are log transformed



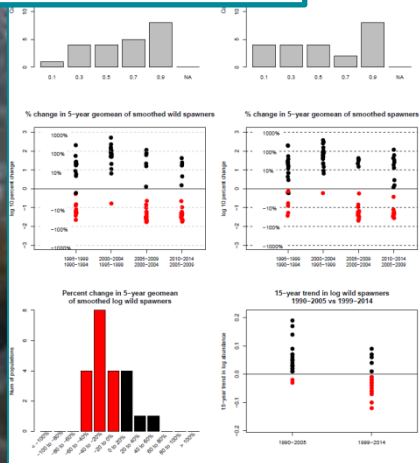
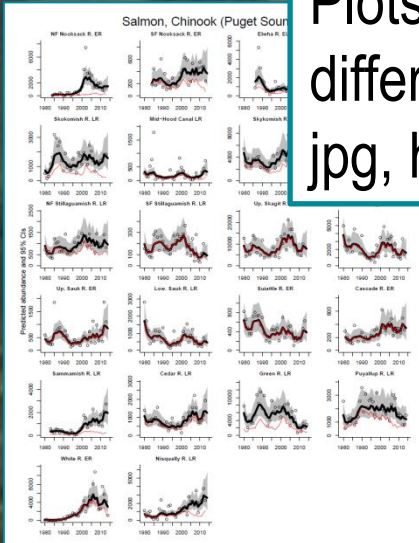
R-based Automated Report (plots/tables) generation

Example from a 5-year status review update

Database (raw data)



Plots and tables in different formats: jpg, html, pdf, ...



```
60
61 ~~~{r summary_fig, echo=FALSE,fig.width=8.5,fig.height=11}-
62 par(mfrow=c(3,2))
63 source("fracwild_tables.R")
64 total.spawners=matdat.spawners[esus==esuname,,drop=FALSE]
65 wild.spawners=matdat.wildspawners[esus==esuname,,drop=FALSE]
66 a=fracwild_table(wild.spawners[pop.to.plot,,drop=FALSE],total.spawners[pop.to.plot,,drop=FALSE])
67
68 barplot(table(cut(as.numeric(as.character(a[,5]))),seq(0,100,by=10)),names.arg=c(as.character(seq(0.1,.9,0.2)),"NA"),ylim=c(0,dim(a)[1]))
69
70 title("Fraction wild across populations 2005-2009")
71
72
73 barplot(table(cut(as.numeric(as.character(a[,6]))),seq(0,100,by=10)),names.arg=c(as.character(seq(0.1,.9,0.2)),"NA"),ylim=c(0,dim(a)[1]))
74
75 title("Fraction wild across populations 2010-2014")
76
77
78 source("geomean_tables.R")
79 a=geomean_table(pop.to.plot,mpg.to.plot,ifit.total,ifit.wild)
80 vals1=vals2=c()
81 geo.start=which(names(a)==%MPG)+1
82 geo.end=which(names(a)==%Change)-1
83 for(i in 1:dim(a)[1]){
84 vals1=rbind(vals1,as.numeric(apply(a[i,geo.start:geo.end],MARGIN=2,FUN=function(x){log10(x)},[1]))))
85 vals2=rbind(vals2,as.numeric(apply(a[i,geo.start:geo.end],MARGIN=2,FUN=function(x){log10(x)},[2]))))
86 vals=sapply(vals,function(x){str_split(x,"D")[[1]][1]})
87 vals2=rbind(vals2,as.numeric(vals))
88
89
90 plot(0:5,0:5,xlim=c(0,4.5),ylim=c(-1*log10(1000),log10(1000)),type="n",axes=F)
91 ylab="log 10 percent change",xlab=""
92 for(i in 1:(dim(vals1)[2]-1)){
93 n=dim(vals2)[1]
94 vals=100*(vals1[,i+1]-vals1[,i])/vals1[,i]-
95 points(i+norm(n,0,0.01),ifelse(vals<0,-1,1)*log10(abs(vals)),
96 cex=1.5,pch=19,col=ifelse(vals<0,"red","black"))
97
98
99 axis(side=1,at=1:4,label=c("1995-1999","1999-1994"))
```

R code embedded in a report generation framework (knitr/Sweave)

Population	MPG	1990-2005	1999-2014
NF Nooksack R. ER	Strait of Georgia	0.07 (0.04, 0.09)	0.04 (0, 0.07)
SF Nooksack R. ER	Strait of Georgia	0.03 (0, 0.06)	-0.06 (-0.1, -0.02)
Elwha R. ER	SJF	-0.02 (-0.06, 0.02)	-0.06 (-0.1, -0.03)
Dungeness R. SR	SJF	0.14 (0.08, 0.19)	0.09 (0.03, 0.14)
Skokomish R. LR	Hood Canal	0.02 (-0.01, 0.05)	-0.07 (-0.11, -0.02)
Mid-Hood Canal LR	Hood Canal	0.03 (0, 0.07)	-0.07 (-0.11, -0.02)
Skykomish R. LR	Whidbey Basin	0.03 (0, 0.06)	-0.02 (-0.04, 0.01)
Snoqualmie R. LR	Whidbey Basin	0.09 (0.05, 0.12)	-0.05 (-0.08, -0.03)
NF Stillaguamish R. LR	Whidbey Basin	0.04 (0.02, 0.06)	-0.04 (-0.06, -0.01)
SF Stillaguamish R. LR	Whidbey Basin	0.01 (-0.01, 0.03)	-0.1 (-0.12, -0.08)
Up. Skagit R. LR	Whidbey Basin	0.07 (0.05, 0.09)	-0.03 (-0.06, 0)
Low. Skagit R. LR	Whidbey Basin	0.05 (0.02, 0.09)	-0.03 (-0.06, -0.01)
Up. Sank R. ER	Whidbey Basin	0.01 (-0.02, 0.04)	0.06 (0.04, 0.08)
Low. Sank R. LR	Whidbey Basin	0.05 (0.01, 0.08)	-0.04 (-0.07, -0.01)
Sniattle R. ER	Whidbey Basin	0.01 (-0.01, 0.03)	-0.01 (-0.04, 0.01)
Cascade R. ER	Whidbey Basin	0.06 (0.04, 0.08)	0.01 (-0.01, 0.03)
Sammamish R. LR	Central/South Sound	0.17 (0.11, 0.23)	-0.02 (-0.06, 0.02)
Cedar R. LR	Central/South Sound	0.03 (0, 0.06)	0.07 (0.05, 0.1)
Green R. LR	Central/South Sound	0.02 (-0.02, 0.06)	-0.12 (-0.16, -0.09)
Puyallup R. LR	Central/South Sound	-0.03 (-0.05, -0.02)	-0.06 (-0.08, -0.03)
White R. ER	Central/South Sound	0.19 (0.17, 0.21)	-0.03 (-0.08, 0.01)
Nisqually R. LR	Central/South Sound	0.05 (0.03, 0.06)	-0.01 (-0.05, 0.03)

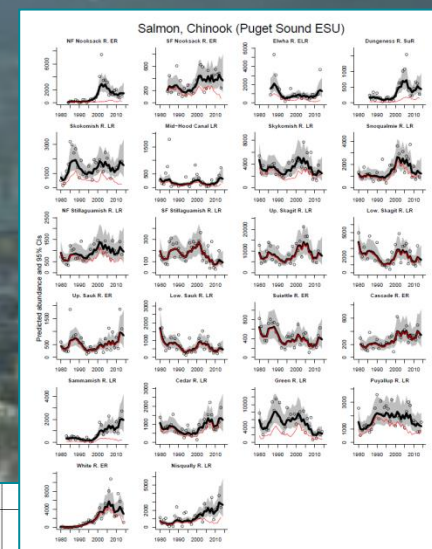
R-based Automated Report (plots/tables) generation

Example from a 5-year status review update

Advantages

- Standardization
- Transparency (“reproducibility”)
- Vastly reduces time to update tables and figures
- Reduces errors
- Easily make analyses available (shiny apps)

6-8 plots
3-5 tables
13 DPS
1-22 pops per DPS



Population	MPG		
NF Nooksack R. ER	Strait of Georgia		
SF Nooksack R. ER	Strait of Georgia		
Elwha R. ELR	SJF	-0.02 (-0.06, 0.02)	-0.06 (-0.1, -0.03)
Dungeness R. SuR	SJF	0.14 (0.08, 0.19)	0.09 (0.03, 0.14)
Skokomish R. LR	Hood Canal	0.02 (-0.01, 0.05)	-0.07 (-0.11, -0.02)
Mid-Hood Canal LR	Hood Canal	0.03 (0, 0.07)	-0.07 (-0.11, -0.02)
Skykomish R. LR	Whidbey Basin	0.03 (0, 0.06)	-0.02 (-0.04, 0.01)
Snoqualmie R. LR	Whidbey Basin	0.09 (0.05, 0.12)	-0.05 (-0.08, -0.03)
NF Stillaguamish R. LR	Whidbey Basin	0.04 (0.02, 0.06)	-0.04 (-0.06, -0.01)
SF Stillaguamish R. LR	Whidbey Basin	0.01 (-0.01, 0.03)	-0.1 (-0.12, -0.08)
Up. Skagit R. LR	Whidbey Basin	0.07 (0.05, 0.09)	-0.03 (-0.06, 0)
Low. Skagit R. LR	Whidbey Basin	0.05 (0.02, 0.09)	-0.03 (-0.06, -0.01)
Up. Sauk R. ER	Whidbey Basin	0.01 (-0.02, 0.04)	0.06 (0.04, 0.08)
Low. Sauk R. LR	Whidbey Basin	0.05 (0.01, 0.08)	-0.04 (-0.07, -0.01)
Suiattle R. ER	Whidbey Basin	0.01 (-0.01, 0.03)	-0.01 (-0.04, 0.01)
Cascade R. ER	Whidbey Basin	0.06 (0.04, 0.08)	0.01 (-0.01, 0.03)
Sammamish R. LR	Central/South Sound	0.17 (0.11, 0.23)	-0.02 (-0.06, 0.02)
Cedar R. LR	Central/South Sound	0.03 (0, 0.06)	0.07 (0.05, 0.1)
Green R. LR	Central/South Sound	0.02 (-0.02, 0.06)	-0.12 (-0.16, -0.09)
Puyallup R. LR	Central/South Sound	-0.03 (-0.05, -0.02)	-0.06 (-0.08, -0.03)
White R. ER	Central/South Sound	0.19 (0.17, 0.21)	-0.03 (-0.08, 0.01)
Nisqually R. LR	Central/South Sound	0.05 (0.03, 0.06)	-0.01 (-0.05, 0.03)

Web-based Protected Species Toolbox and Cloud-based Platform for Running Tools

This PDF was later amended to make the document 508 compliant.

- **Project accomplishments:**
 - Content manager website built (meets NOAA IT certifications)
 - <https://www.st.nmfs.noaa.gov/npst>
 - Public R/Shiny server established at NWFSC
 - <https://dataexplorer.northwestscience.fisheries.noaa.gov/>
 - Multiple R shiny applications (agTrend, VRAP, ROSE, Growth, PVAwidget)