# American Samoa Model Checks

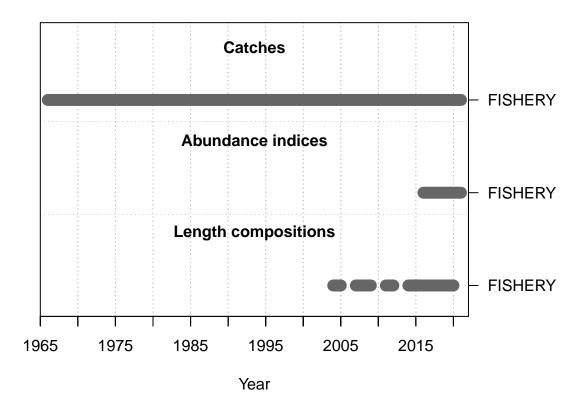
Meg Oshima

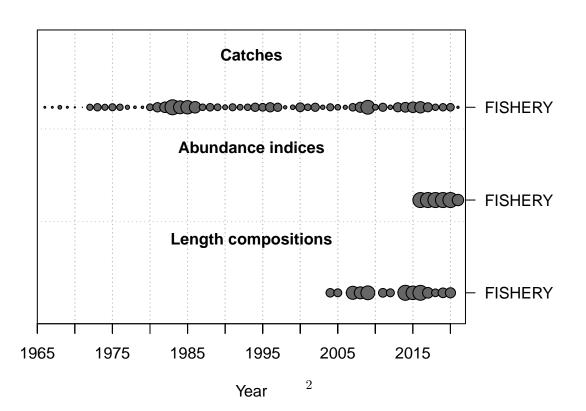
2022-08-16

This is a summary report for the APVI base model run.

### **Model Output**

### Input Data





#### Convergence Check

Converged

```
## 1 TRUE 1.42575e-05

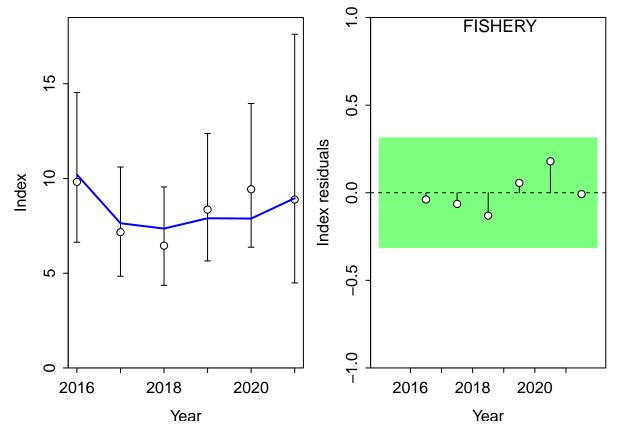
## [1] "1 NOTE: Max data length bin: 85 < max pop len bins: 94; so will accumulate larger pop len bin
## [2] "2 Minimum pop size bin:_1; is > L at Amin for sex: 1; Gpat: 1; L= 0"
## [3] "3 warning: poor convergence in Fmsy, final dy/dy2= -0.00793234"
## [4] "N warnings: 3"
```

#### Fit to Model

#### **CPUE**

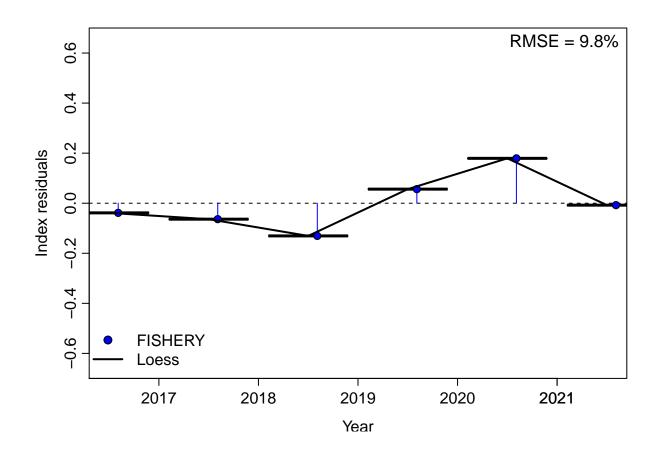
##
## Running Runs Test Diagnosics for Index
## Plotting Residual Runs Tests

MaxGrad



```
##
## Runs Test stats by Index:
## Plotting JABBA residual plot
## Warning in simpleLoess(y, x, w, span, degree = degree, parametric = parametric, : Chernobyl! trL>n 6
```

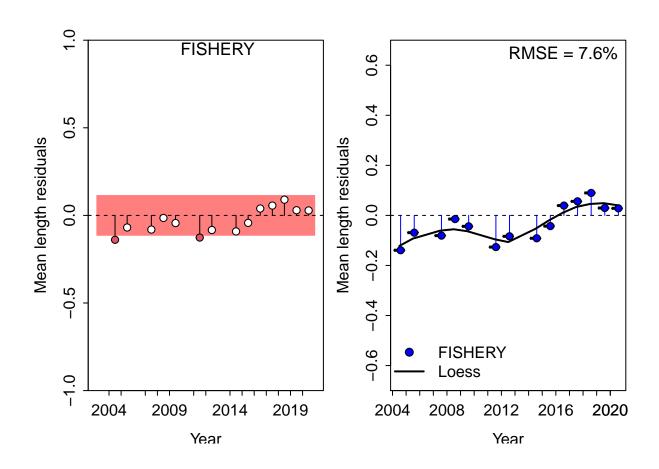
## Warning in simpleLoess(y, x, w, span, degree = degree, parametric = parametric, : Chernobyl! trL>n 6



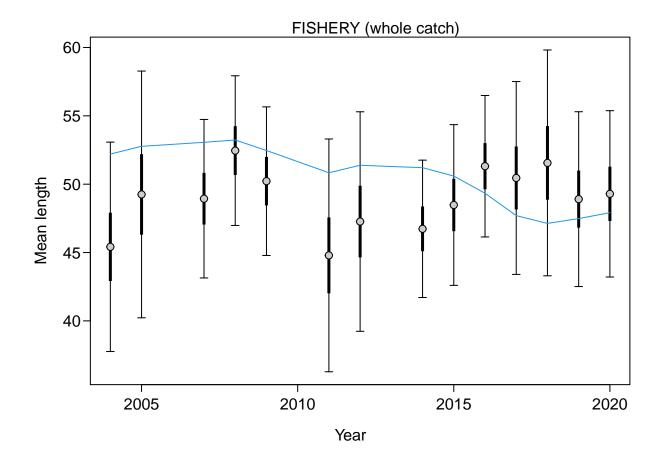
##
## RMSE stats by Index:

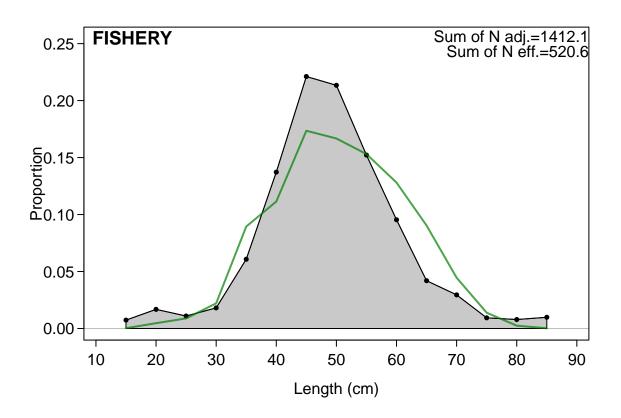
#### Length Comp

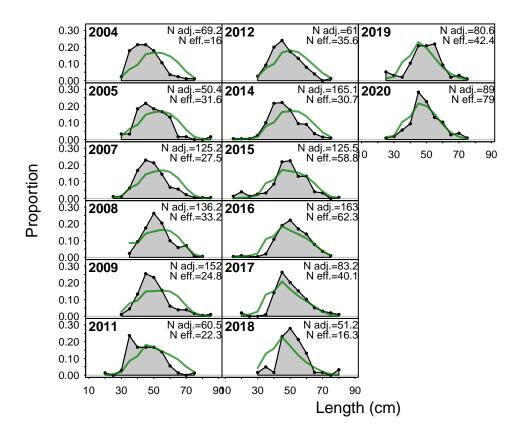
#Factor	Fleet	$New\_Var\_adj$	Type	Name
4	1	0.106439	len	FISHERY



##
## RMSE stats by Index:







#### Retrospective and Hindcasting

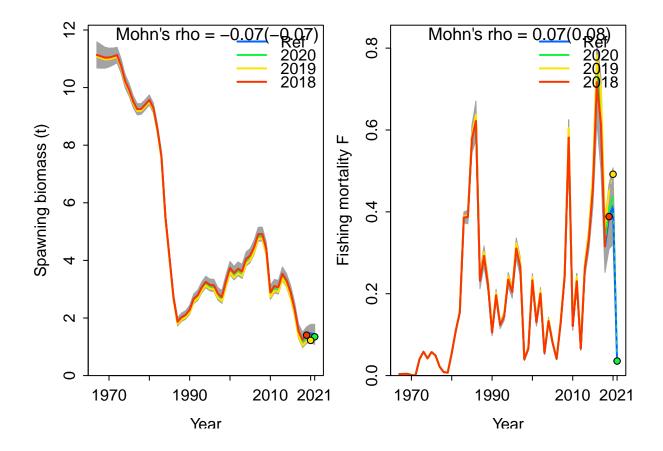
#### Retrospective

## Plotting Retrospective pattern

##

 $\mbox{\tt \#\#}$  Mohn's Rho stats, including one step ahead forecasts:

## Plotting Retrospective pattern



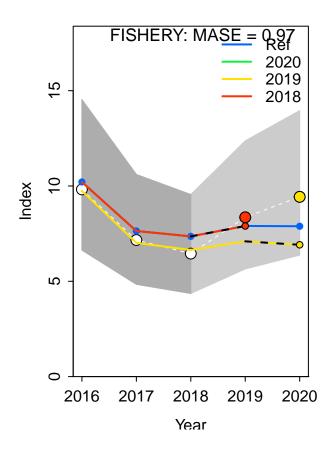
## Mohn's Rho stats, including one step ahead forecasts:

```
## type peel Rho ForecastRho
## 1 F 2020 0.0527939515 0.0529898255
## 2 F 2019 0.1596800272 0.1874511167
## 3 F 2018 0.0001044998 0.0001055591
## 4 F Combined 0.0708594928 0.0801821671
```

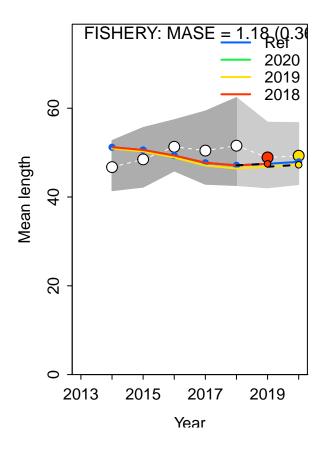
#### Hindcasting

```
## Plotting Hindcast Cross-Validation (one-step-ahead)
##
## Computing MASE with only 2 of 3 prediction residuals for Index FISHERY
##
```

## Warning: Unequal spacing of naive predictions residuals may influence the interpretation of MASE



```
##
## MASE stats by Index:
## Plotting Hindcast Cross-Validation (one-step-ahead)
##
## Computing MASE with only 2 of 3 prediction residuals for Index FISHERY
##
## Warning: Unequal spacing of naive predictions residuals may influence the interpretation of MASE
##
## MASE stats by Index:
```



#### **Recruitment Deviations**

## Skipped SSplotrecdevs - no rec devs estimated

#### Likelihood Profile

```
## [1] "SR_LN"
```

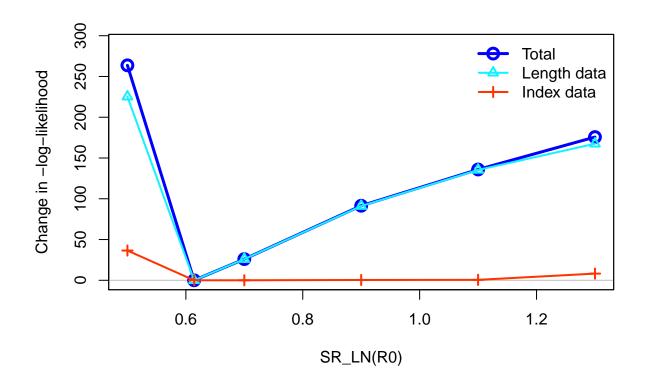
- ## Parameter matching profile.string=SR\_LN: SR\_LN(R0)
- ## Parameter values (after subsetting based on input 'models'): 0.5, 0.7, 0.9, 1.1, 1.3, 0.614098

##

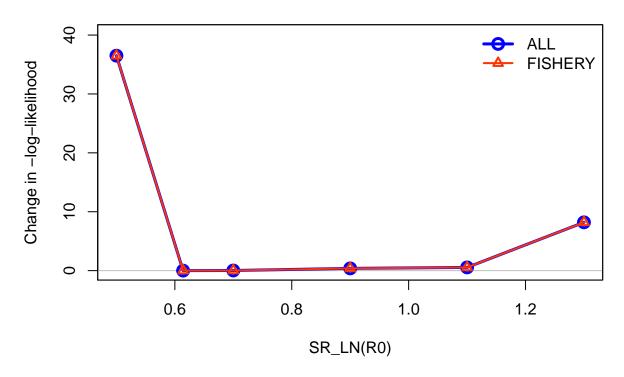
- ## Likelihood components showing max change as fraction of total change.
- ## To change which components are included, change input 'minfraction'.

##		frac_change	include	label
##	TOTAL	1.0000	TRUE	Total
##	Catch	0.0021	FALSE	Catch
##	Equil_catch	0.0053	FALSE	Equilibrium catch
##	Survey	0.1384	TRUE	Index data
##	Length_comp	0.8541	TRUE	Length data
##	Recruitment	0.0000	FALSE	Recruitment

```
## InitEQ_Regime
                             0.0000
                                      FALSE Initital equilibrium recruitment
## Forecast_Recruitment
                             0.0000
                                      FALSE
                                                         Forecast recruitment
## Parm_priors
                             0.0000
                                      FALSE
                                                                       Priors
## Parm_softbounds
                             0.0000
                                      FALSE
                                                                  Soft bounds
## Parm_devs
                             0.0000
                                      FALSE
                                                         Parameter deviations
## Crash_Pen
                             0.0000
                                      FALSE
                                                                Crash penalty
## Parameter matching profile.string = 'SR_LN': 'SR_LN(RO)
## Parameter values (after subsetting based on input 'models'): 0.5, 0.7, 0.9, 1.1, 1.3, 0.614098,
## Fleet-specific likelihoods showing max change as fraction of total change.
## To change which components are included, change input 'minfraction'.
                         frac_change include
## prof.table....c.1.3..
                                   1
                                        TRUE
```

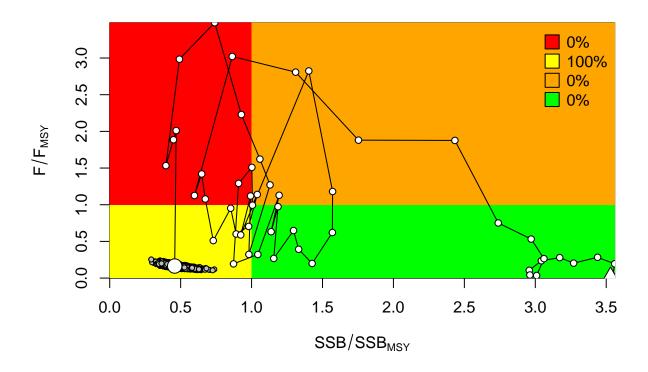


# Changes in survey likelihood by fleet

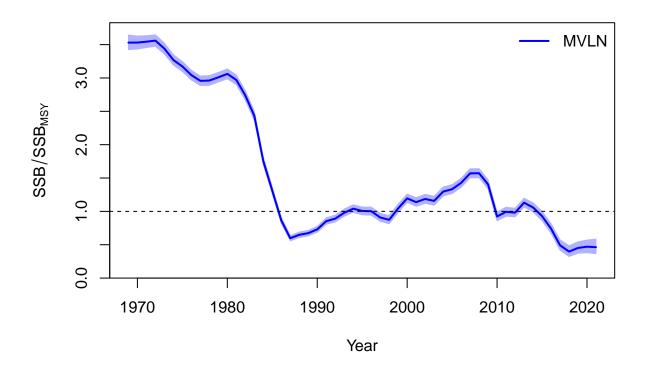


### Management Quantities

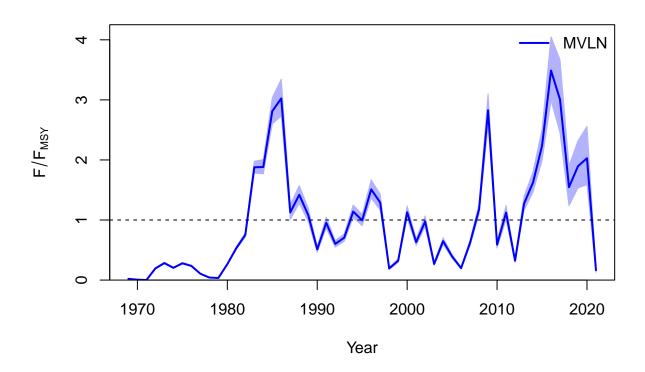
```
##
## starter.sso with Bratio: SSB/SSBMSY and F: _abs_F
##
```



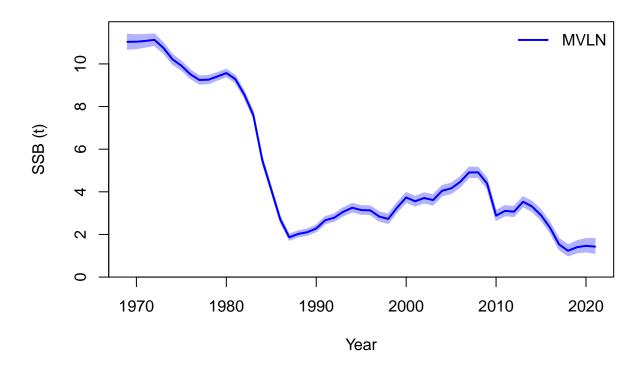
##
## Plot Comparison of stock



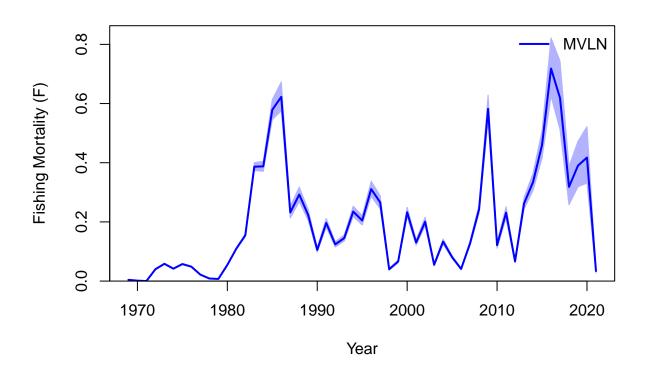
## Plot Comparison of harvest



## Plot Comparison of SSB



## Plot Comparison of F



## RStudioGD ## 2

# Jitter

