American Samoa Model Checks

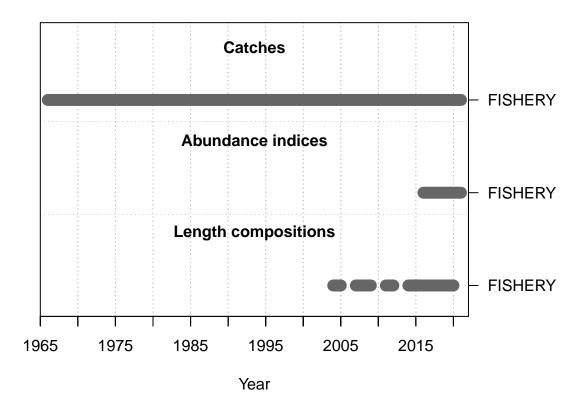
Meg Oshima

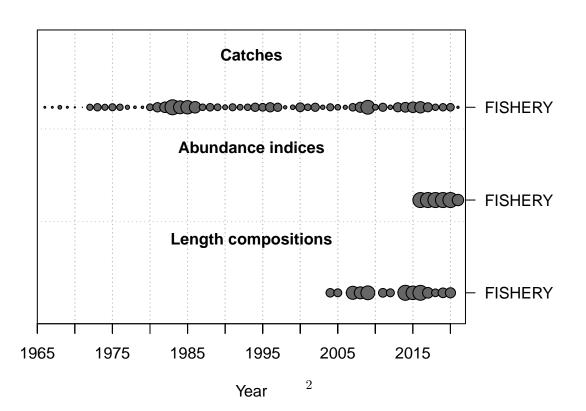
2022-08-11

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

Model Output

Input Data





Convergence Check

Converged

```
## 1 TRUE 6.80005e-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= -0.00555695"

## [4] " N parameters are on or within 1% of min-max bound: 2; check results, variance may be suspect"

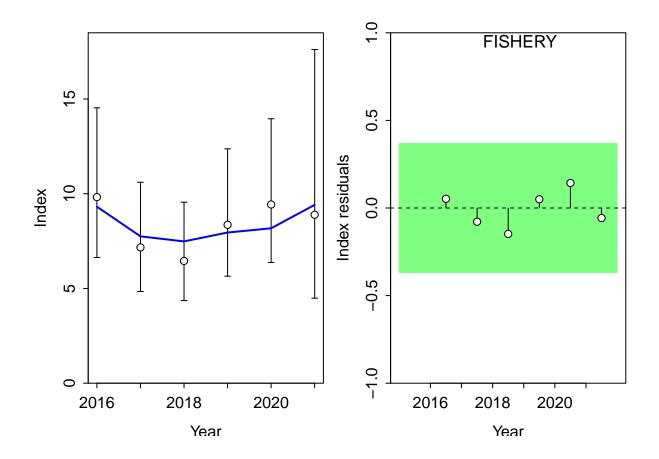
## [5] "N warnings: 3"
```

Fit to Model

CPUE

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

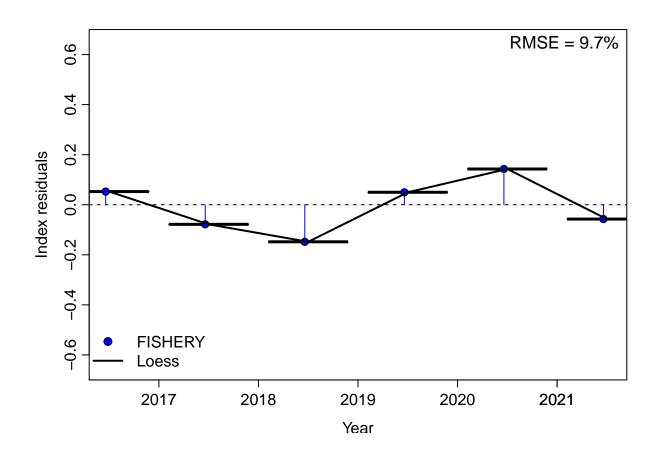
MaxGrad



```
## 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
Warning in sqrt(sum.squares/one.delta): NaNs produced



RMSE stats by Index:

Length Comp

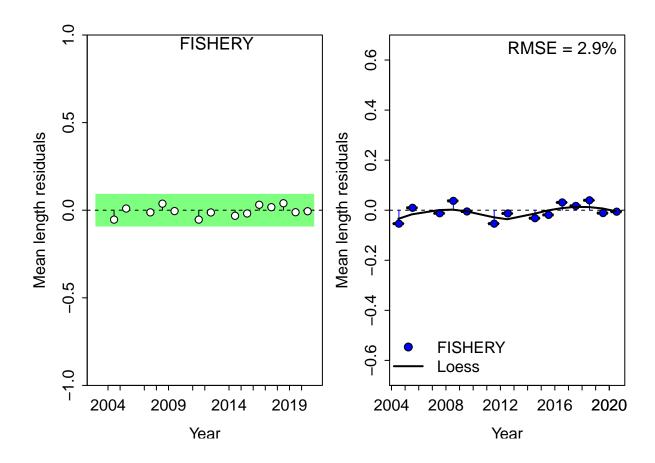
$\# {\operatorname{Factor}}$	Fleet	New_Var_adj	Type	Name
4	1	0.461202	len	FISHERY

##
Running Runs Test Diagnosics for Mean length
Plotting Residual Runs Tests
##

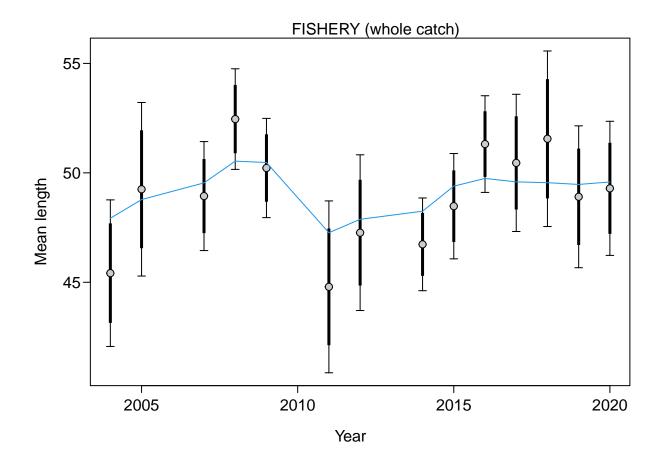
Runs Test stats by Mean length:

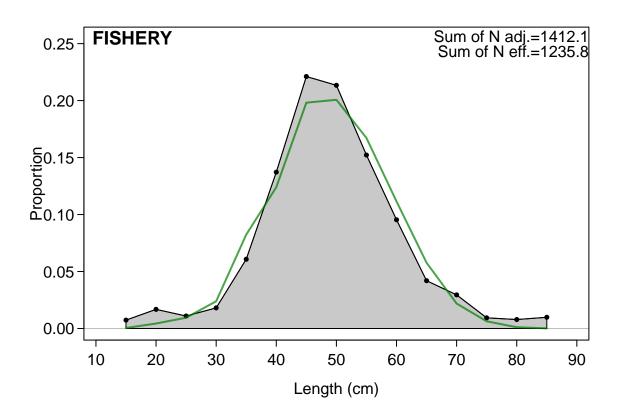
Index runs.p test sigma3.lo sigma3.hi type ## 1 FISHERY 0.397 Passed -0.09035894 0.09035894 len

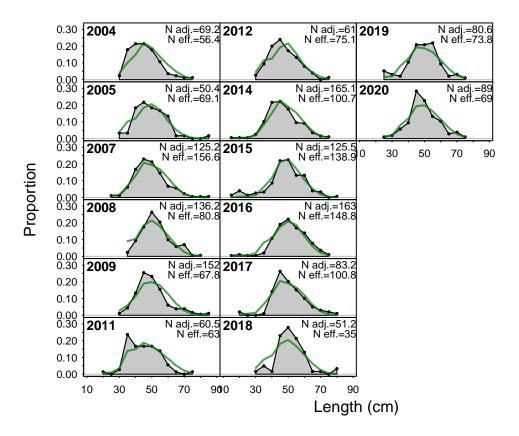
Plotting JABBA residual plot



##
RMSE stats by Index:







Retrospective and Hindcasting

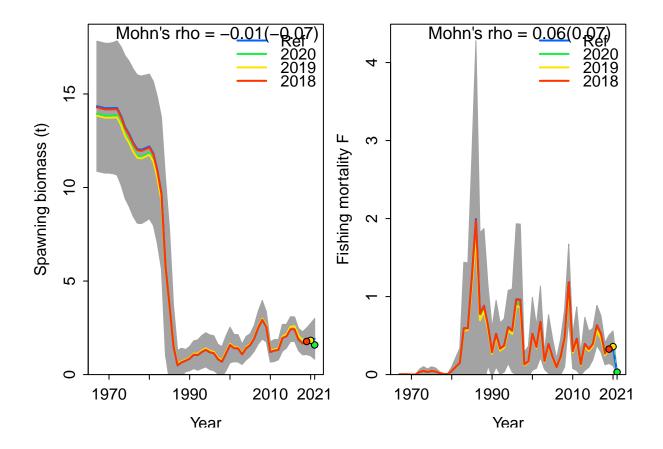
Retrospective

Plotting Retrospective pattern

##

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

Plotting Retrospective pattern



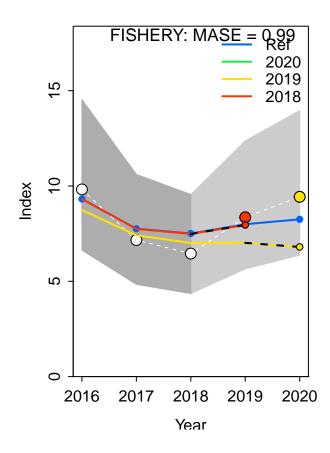
Mohn's Rho stats, including one step ahead forecasts:

```
## type peel Rho ForecastRho
## 1 F 2020 0.177554055 0.135189424
## 2 F 2019 0.008035836 0.066389079
## 3 F 2018 0.005667631 0.009267606
## 4 F Combined 0.063752507 0.070282036
```

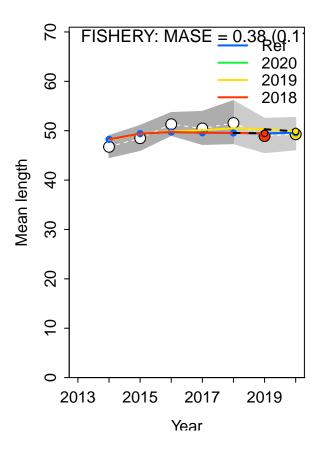
Hindcasting

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

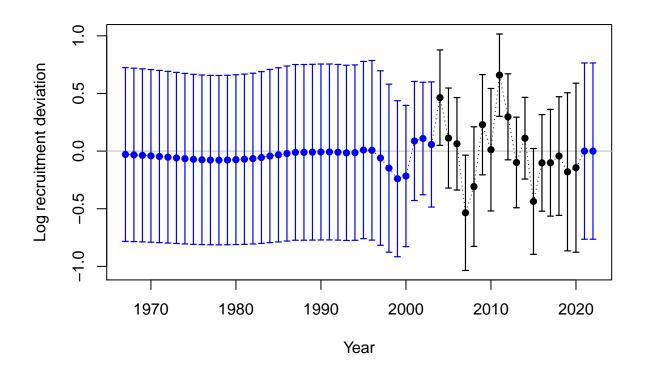
 $\hbox{\tt\#\#-Warning:}\quad \hbox{\tt Unequal spacing of naive predictions residuals may influence the interpretation of MASE}$



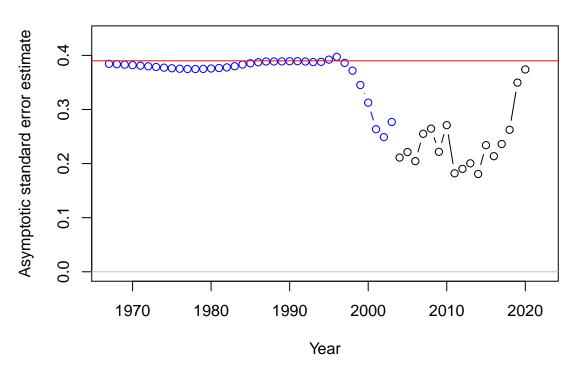
```
##
## MASE stats by Index:
## Plotting Hindcast Cross-Validation (one-step-ahead)
##
## Computing MASE with only 2 of 3 prediction residuals for Index FISHERY
##
## Warning: Unequal spacing of naive predictions residuals may influence the interpretation of MASE
##
## MASE stats by Index:
```



Recruitment Deviations

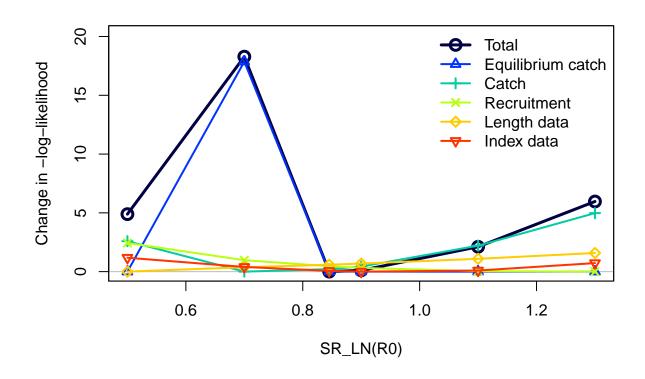


Recruitment deviation variance

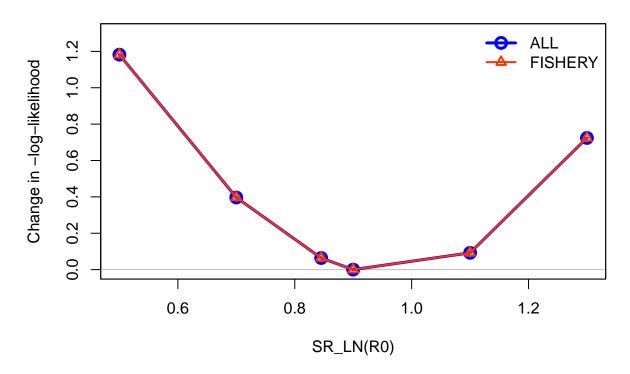


Likelihood Profile

```
## [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, 0.845241
## 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
                                                                       Total
## Catch
                            0.2721
                                      TRUE
                                                                      Catch
## Equil_catch
                            0.9769
                                      TRUE
                                                           Equilibrium catch
## Survey
                            0.0646
                                      TRUE
                                                                 Index data
## Length comp
                            0.0866
                                      TRUE
                                                                Length data
## Recruitment
                            0.1332
                                    TRUE
                                                                Recruitment
## InitEQ_Regime
                            0.0000 FALSE Initital equilibrium recruitment
## Forecast_Recruitment
                          0.0000 FALSE
                                                       Forecast recruitment
## Parm_priors
                            0.0000 FALSE
                                                                     Priors
                            0.0000 FALSE
                                                                Soft bounds
## Parm_softbounds
                                   FALSE
## Parm devs
                            0.0000
                                                       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, 0.845241,
## 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..
```

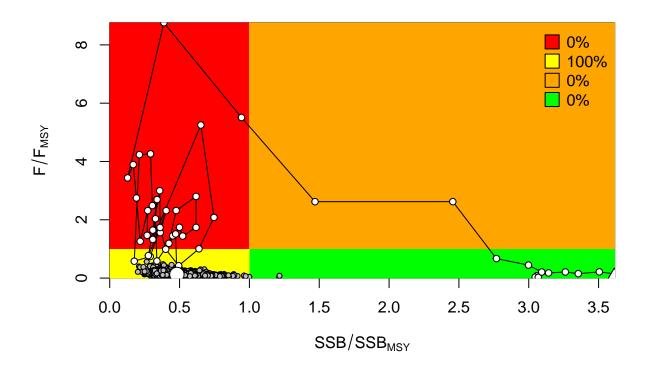


Changes in survey likelihood by fleet

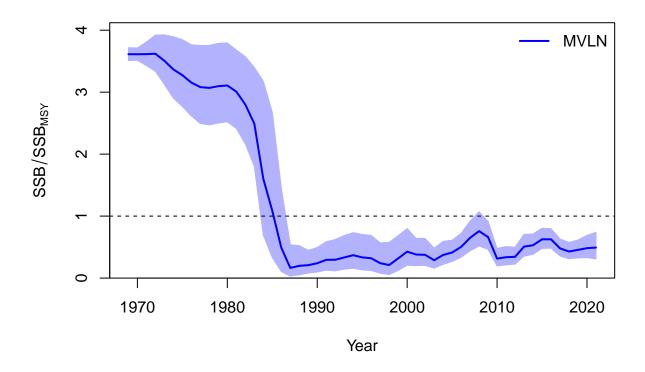


Management Quantities

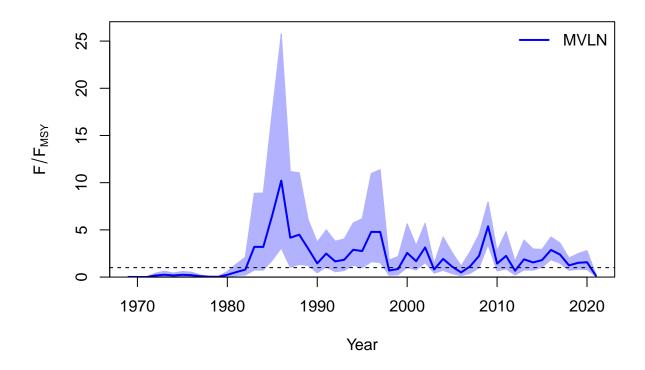
```
##
## starter.sso with Bratio: SSB/SSBMSY and F: _abs_F
##
```



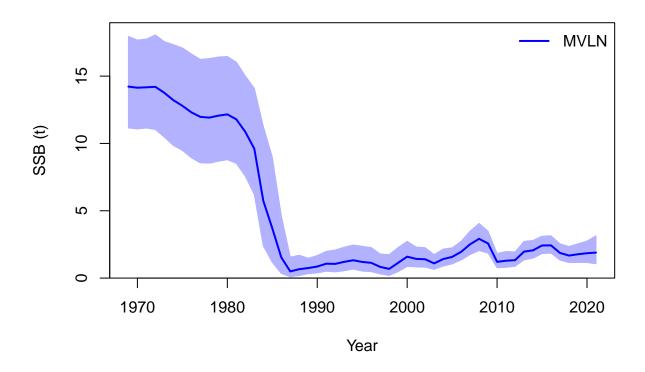
##
Plot Comparison of stock



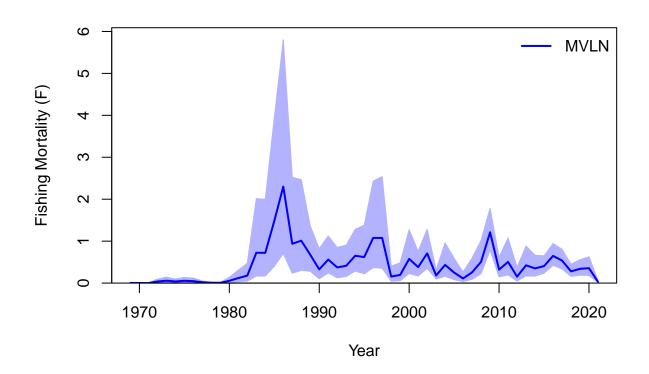
Plot Comparison of harvest



##
Plot Comparison of SSB



Plot Comparison of F



RStudioGD ## 2

Jitter

