American Samoa Model Checks

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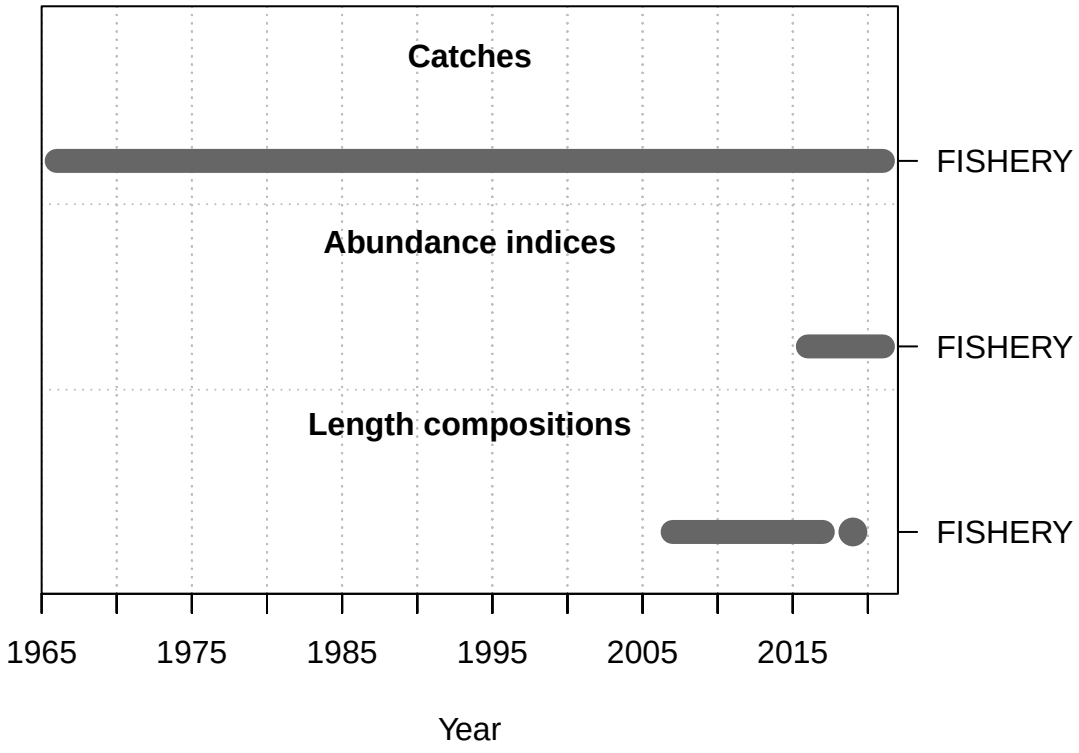
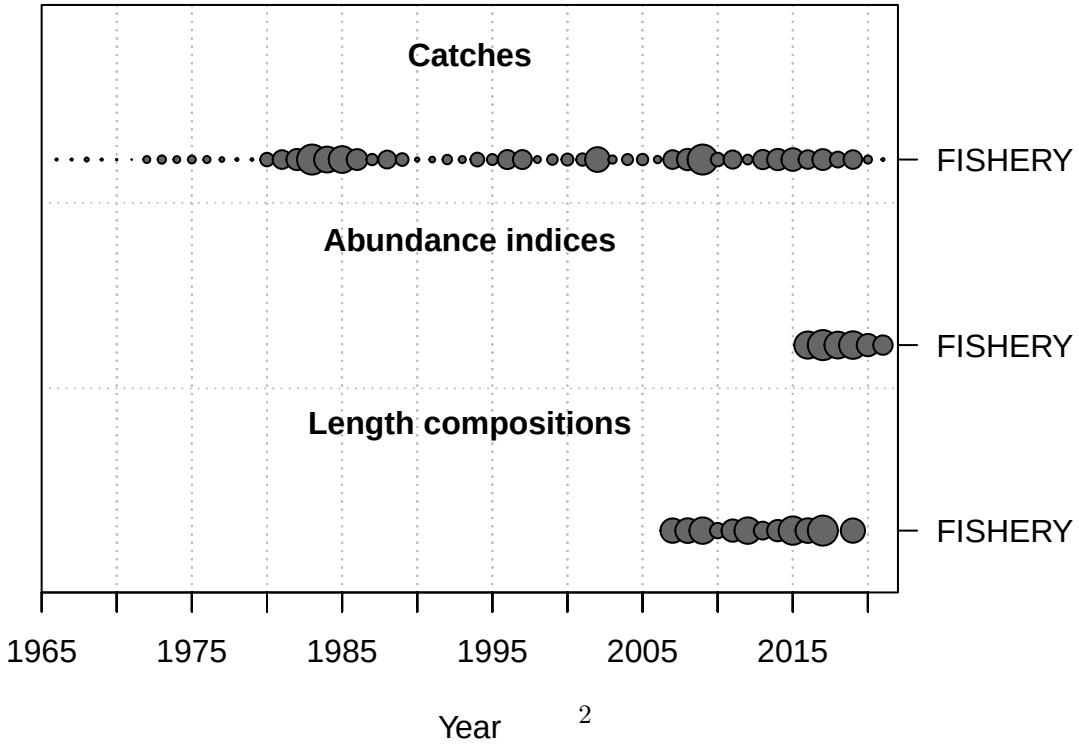
This is a summary report for the APRU base model run.

Model Output

Input Data





Convergence Check

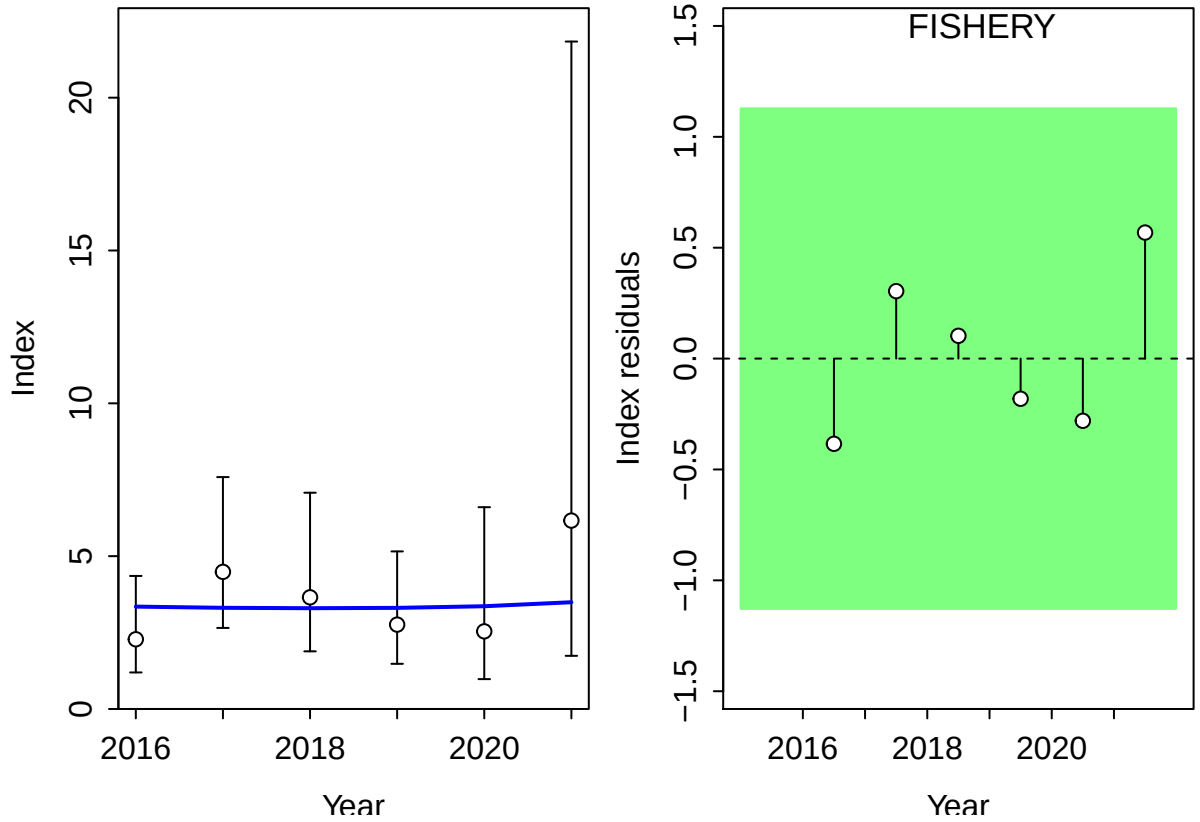
```
##   Converged     MaxGrad
## 1      TRUE 3.39772e-05
```

```
## [1] "1 NOTE:  Max data length bin: 90  < max pop len bins: 100; so will accumulate larger pop len bi
## [3] "N warnings: 1"
```

Fit to Model

CPUE

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

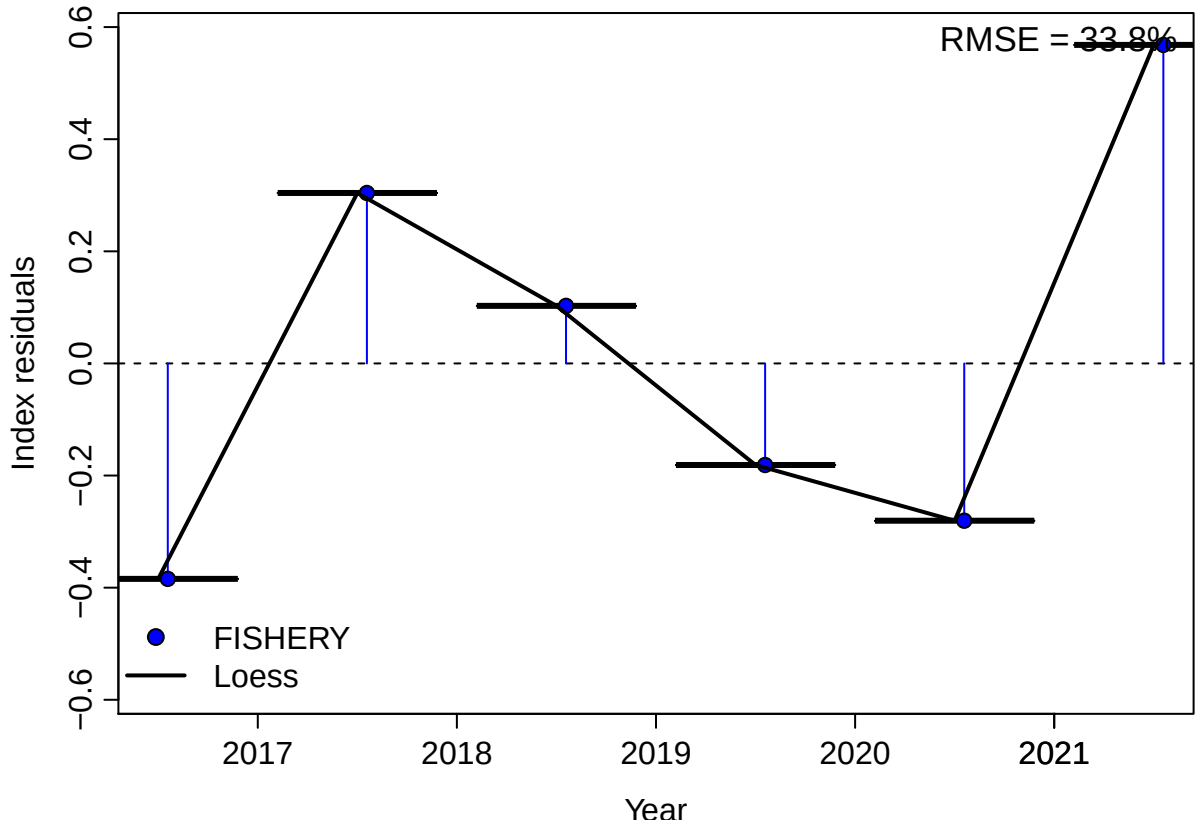



```
##
## Runs Test stats by Index:
## Plotting JABBA residual plot
```

```
## 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
```

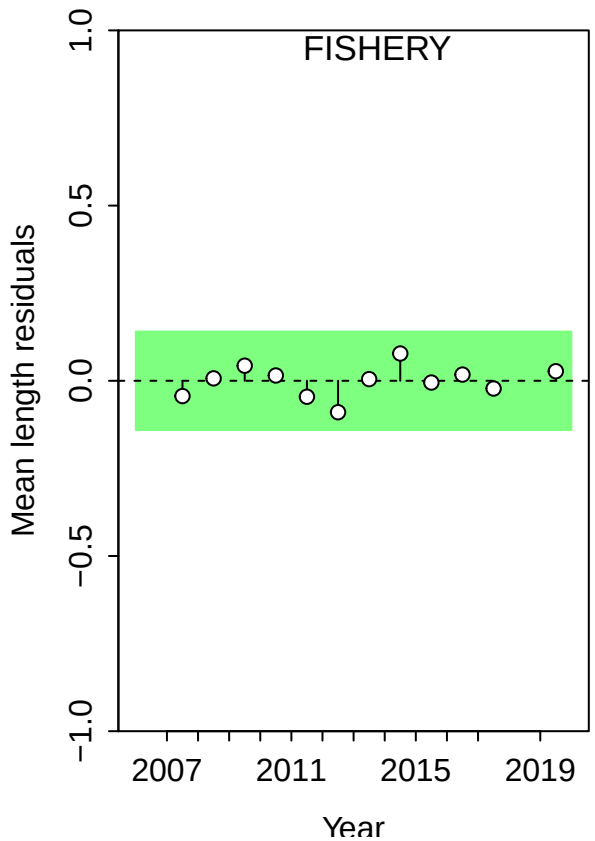


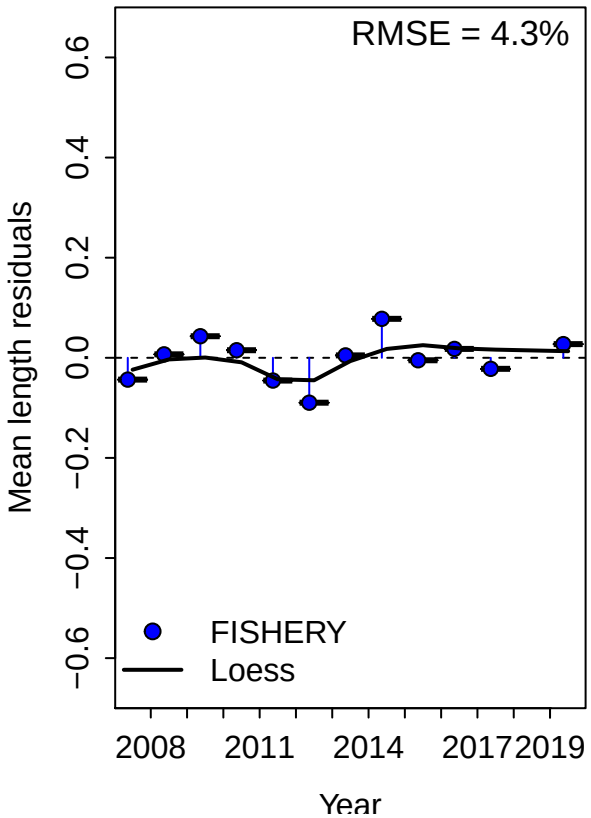

```
##
## RMSE stats by Index:
```

Length Comp

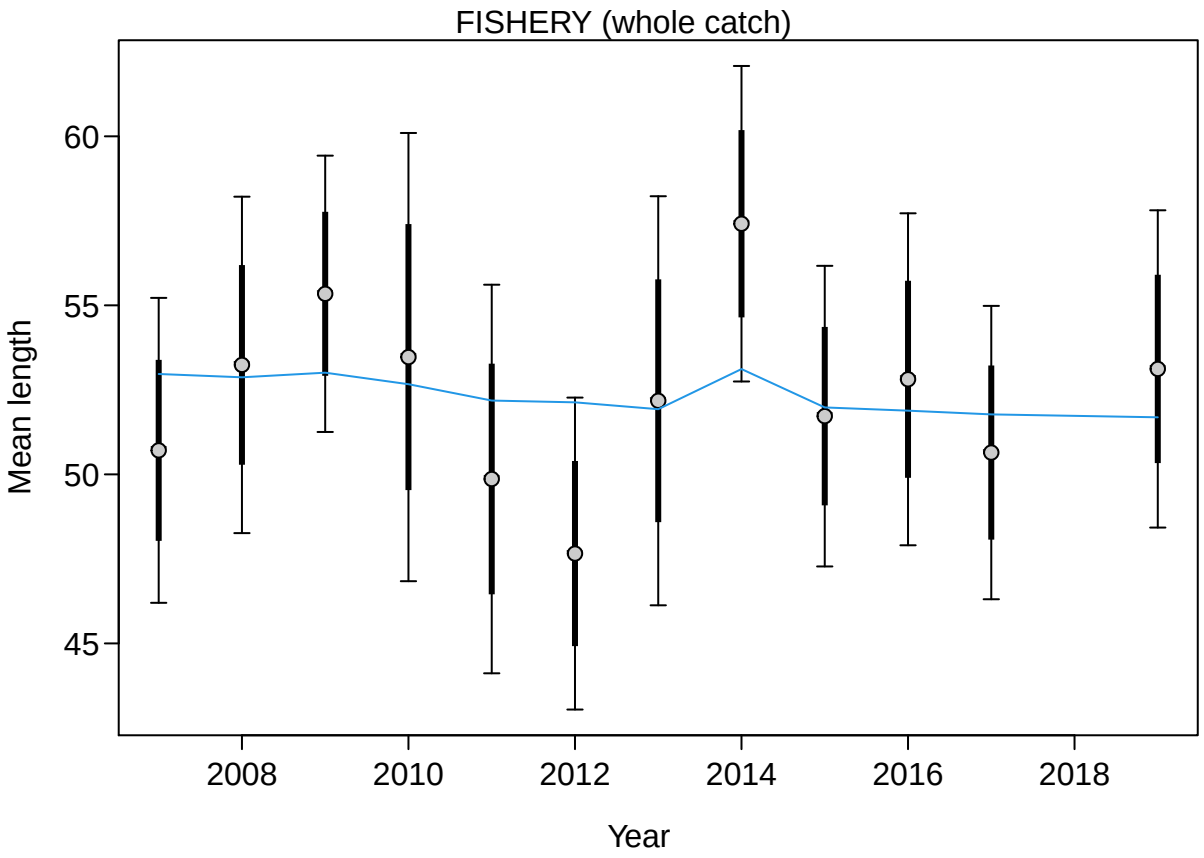
#Factor	Fleet	New_Var_adj	Type	Name
4	1	0.352304	len	FISHERY

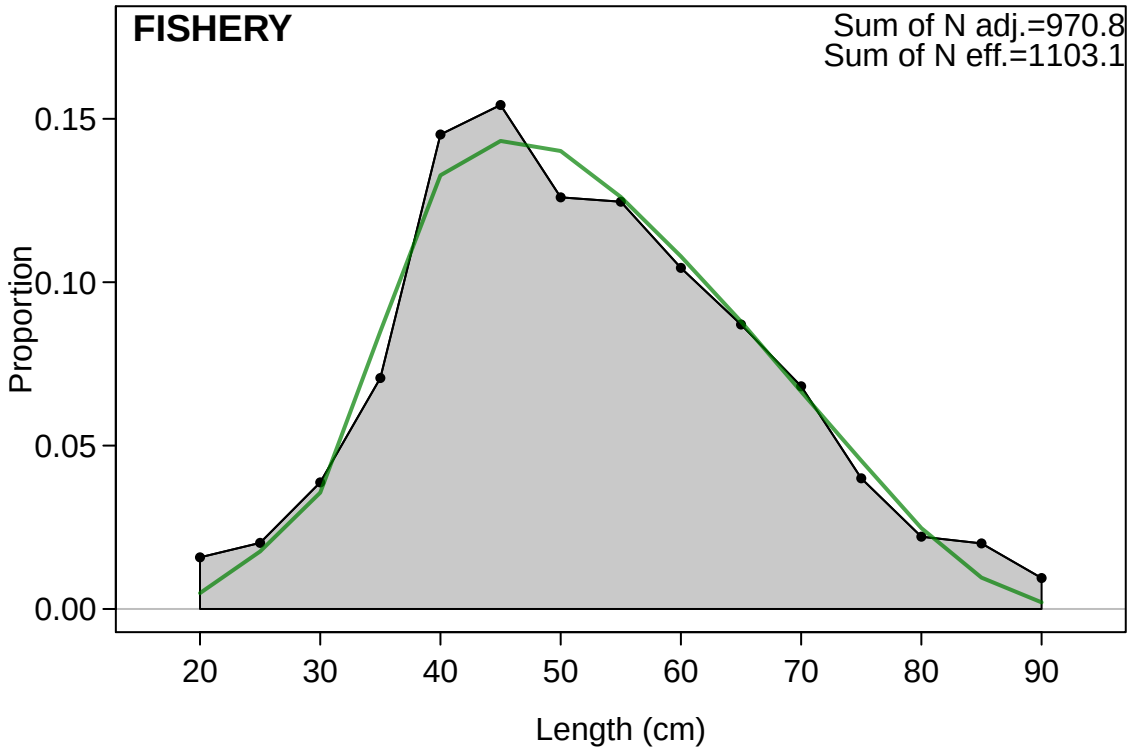
```
##
##  Running Runs Test Diagnosics for Mean length
## Plotting Residual Runs Tests

##
## Runs Test stats by Mean length:

##     Index runs.p   test  sigma3.lo sigma3.hi type
## 1 FISHERY  0.767 Passed -0.1406156 0.1406156  len

## Plotting JABBA residual plot
```

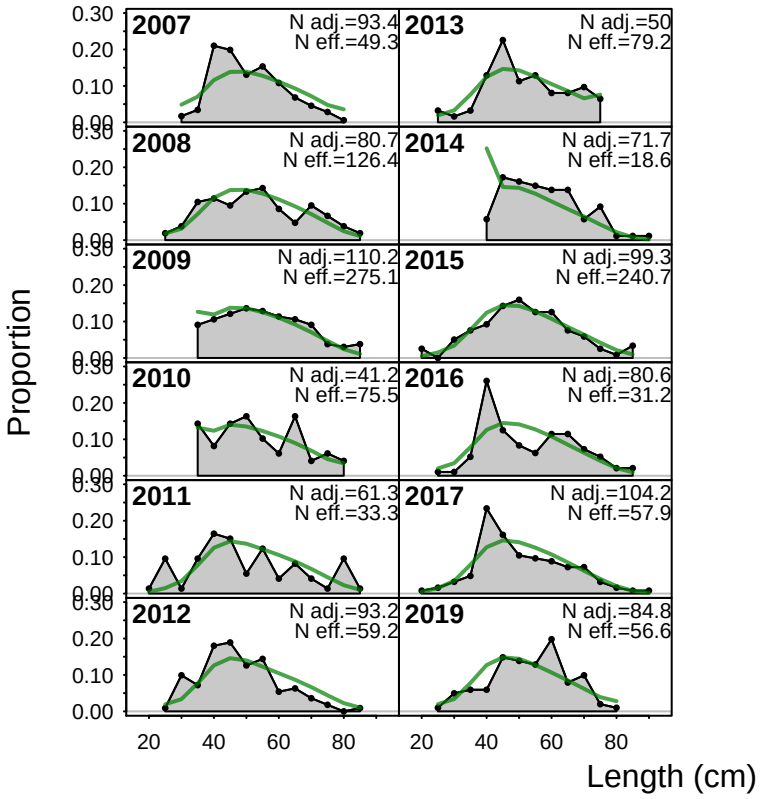





```
##
## RMSE stats by Index:

##    indices RMSE.perc nobs
## 1  FISHERY       4.3   12
## 2 Combined       4.3   12
```


FISHERY (whole catch)
Mean length
60
55
50
45
2008
2010
2012
2014
2016
2018
Year


FISHERY
Sum of N adj.=970.8
Sum of N eff.=1103.1
Proportion
0.15
0.10
0.05
0.00
20
30
40
50
60
70
80
90
Length (cm)

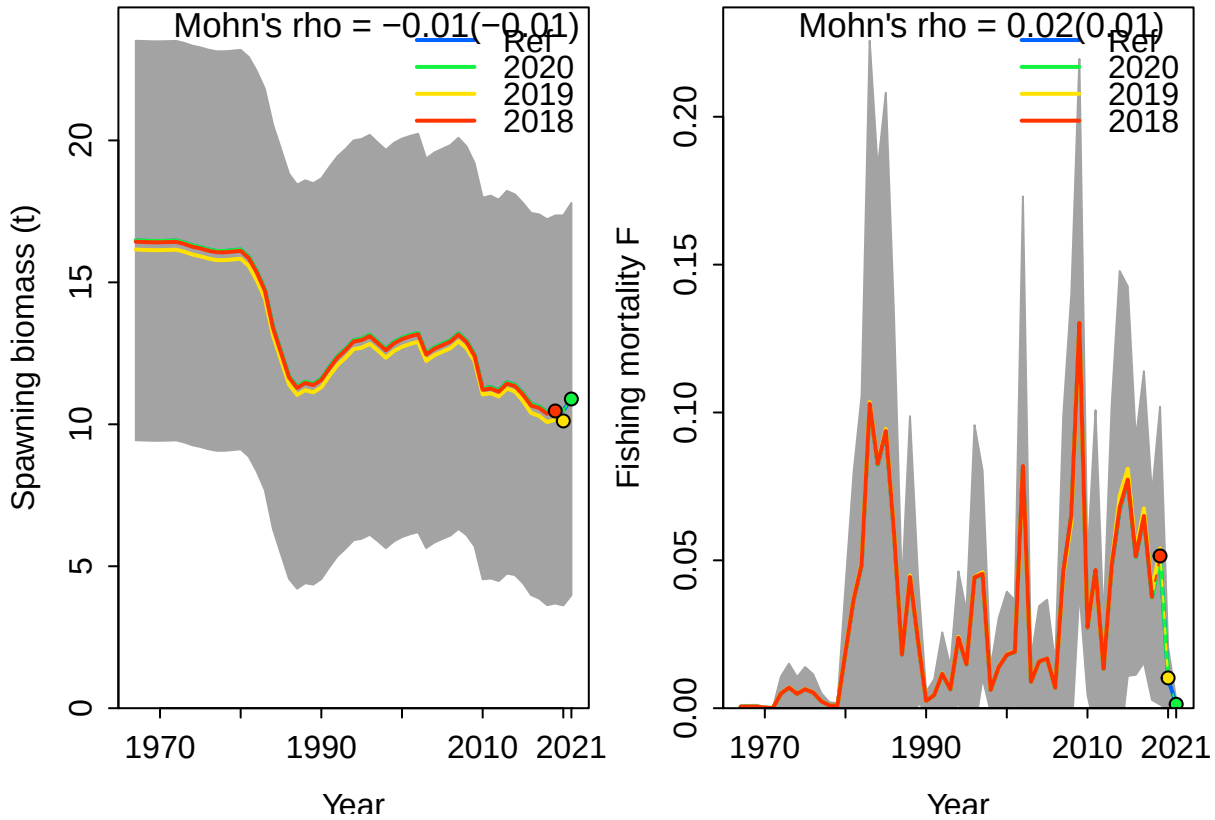


Retrospective and Hindcasting

Retrospective

```
## Plotting Retrospective pattern
```

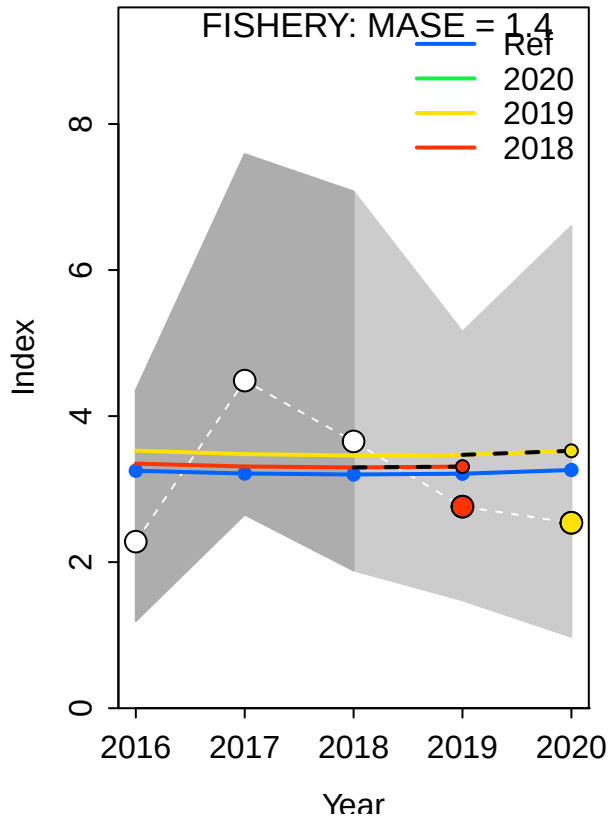
```
##
## Mohn's Rho stats, including one step ahead forecasts:
## Plotting Retrospective pattern
```




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

##   type     peel          Rho  ForecastRho
## 1    F     2020 -0.001051887 -0.000178428
## 2    F     2019  0.048719589  0.030305410
## 3    F     2018  0.002404493 -0.001775710
## 4    F Combined  0.016690732  0.009450424
```

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)
##
## No observations in evaluation years to compute prediction residuals for Index FISHERY
##
## 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)
```

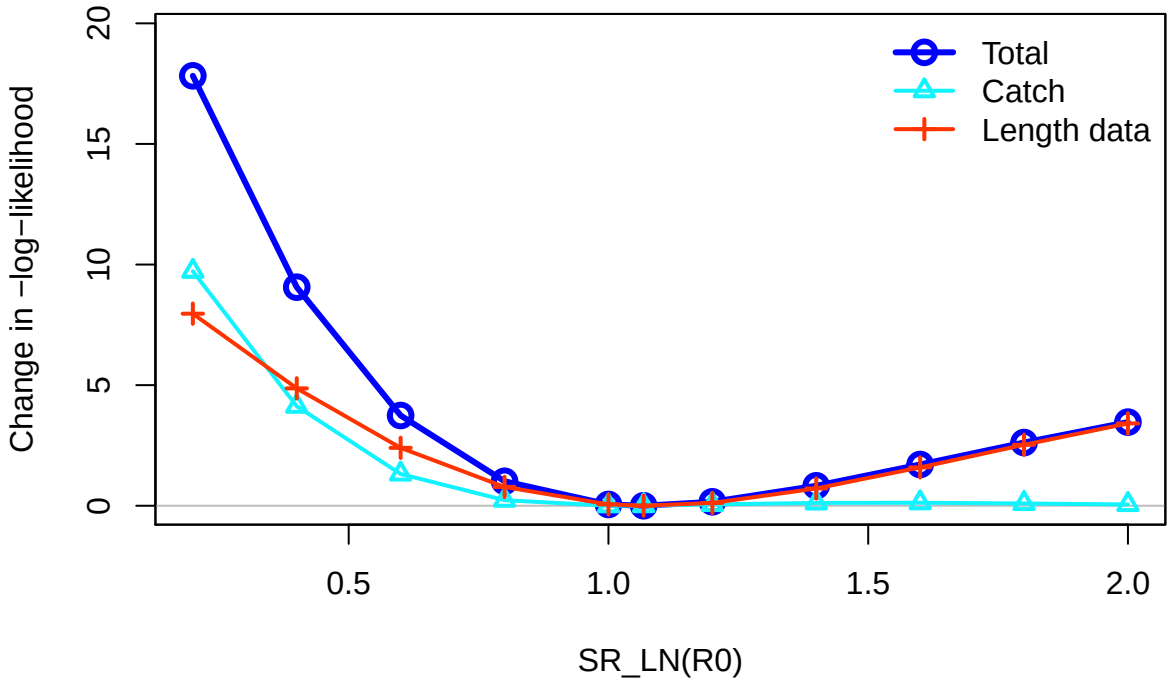
```
## Parameter values (after subsetting based on input 'models'): 0.2, 0.4, 0.6, 0.8, 1, 1.2, 1.4, 1.6, 1
```

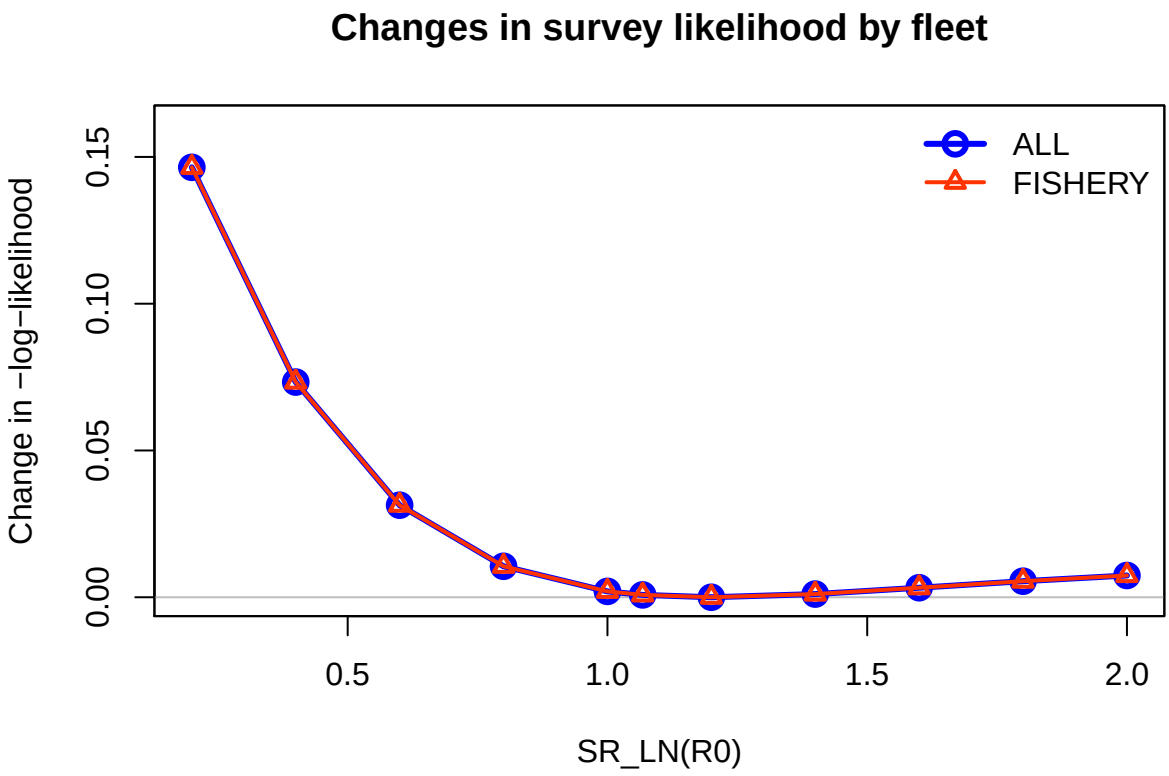
```
##
## Likelihood components showing max change as fraction of total change.
## To change which components are included, change input 'minfraction'.
```

```
##                       frac_change include                           label
## TOTAL                      1.0000    TRUE                           Total
## Catch                      0.5454    TRUE                           Catch
## Equil_catch                0.0000   FALSE                 Equilibrium catch
## Survey                     0.0082   FALSE                       Index data
## Length_comp                0.4467    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(R0)
## Parameter values (after subsetting based on input 'models'): 0.2, 0.4, 0.6, 0.8, 1, 1.2, 1.4, 1.6, 1
```

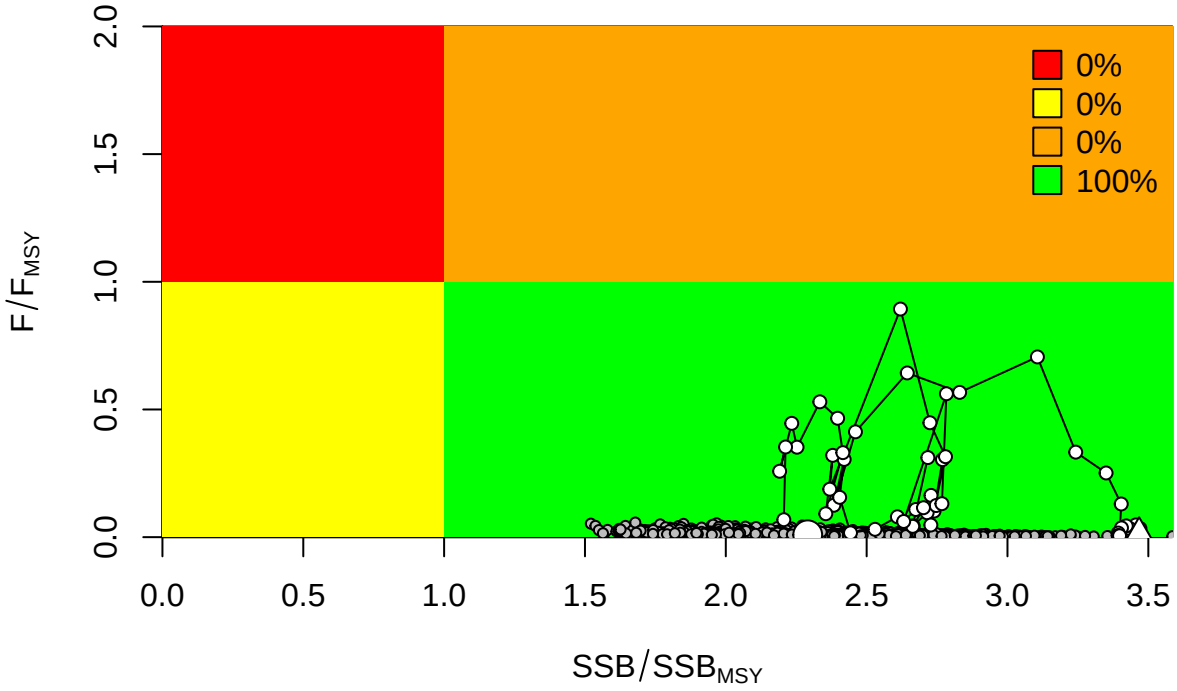
```
## Fleet-specific likelihoods showing max change as fraction of total change.
## To change which components are included, change input 'minfraction'.
##                        frac_change include
## prof.table....c.1.3..            1    TRUE
```



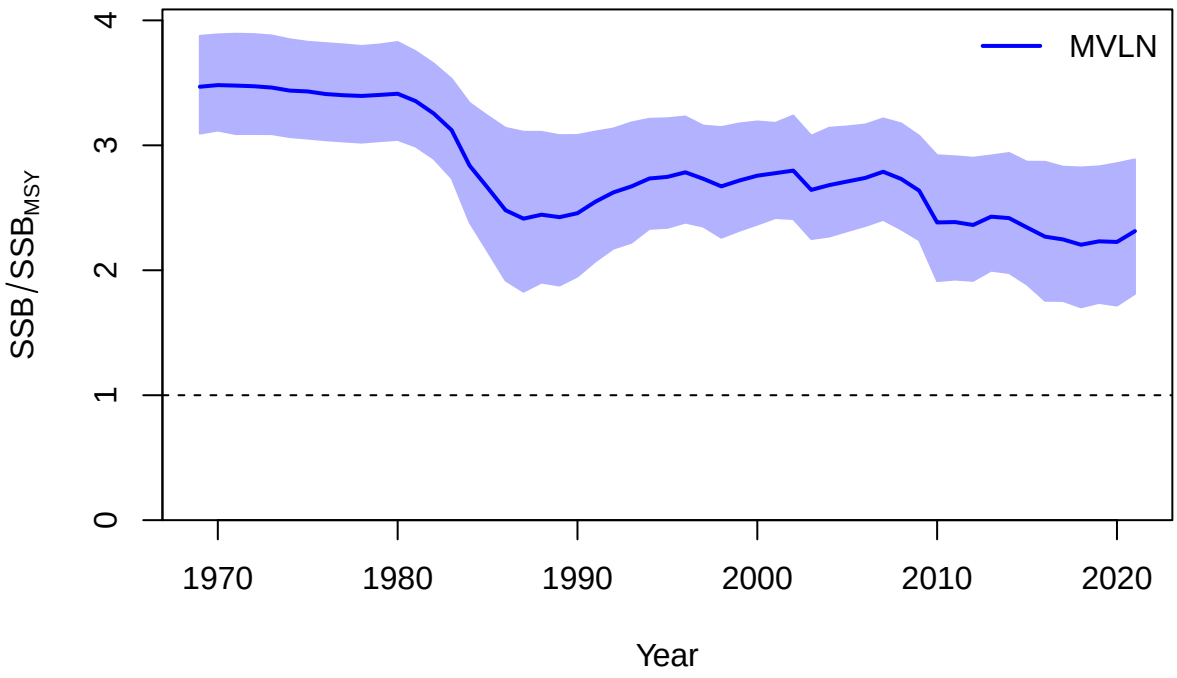


Management Quantities

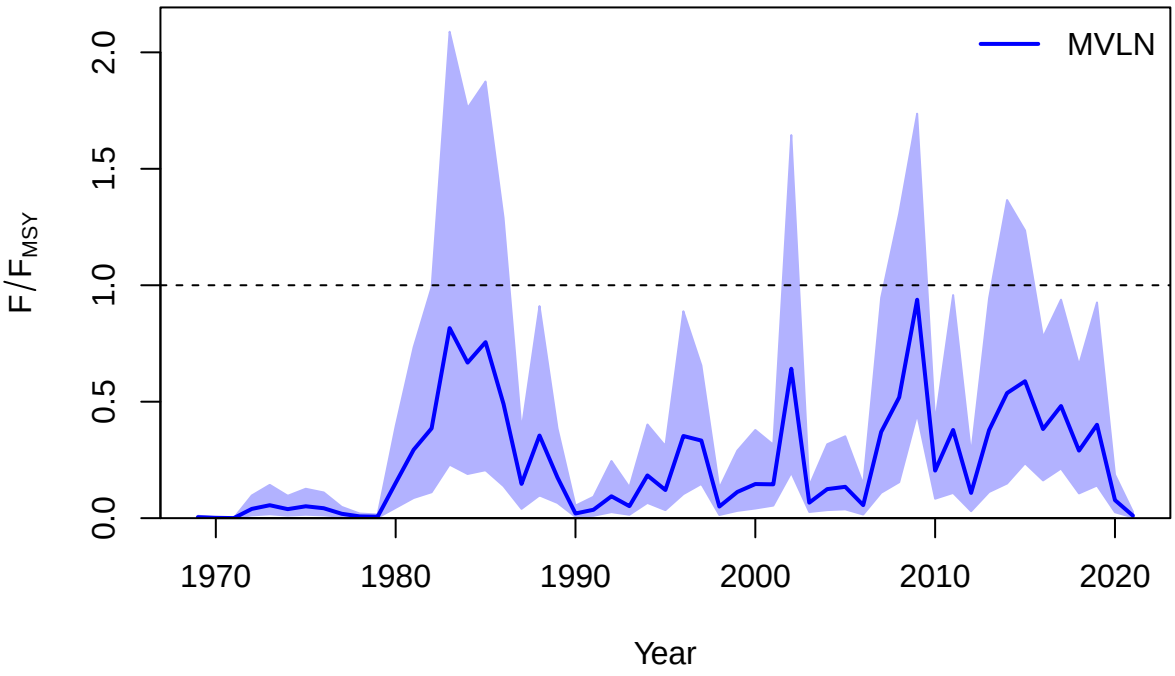
```
##
##  starter.sso with Bratio: SSB/SSBMSY and F: _abs_F
##
```



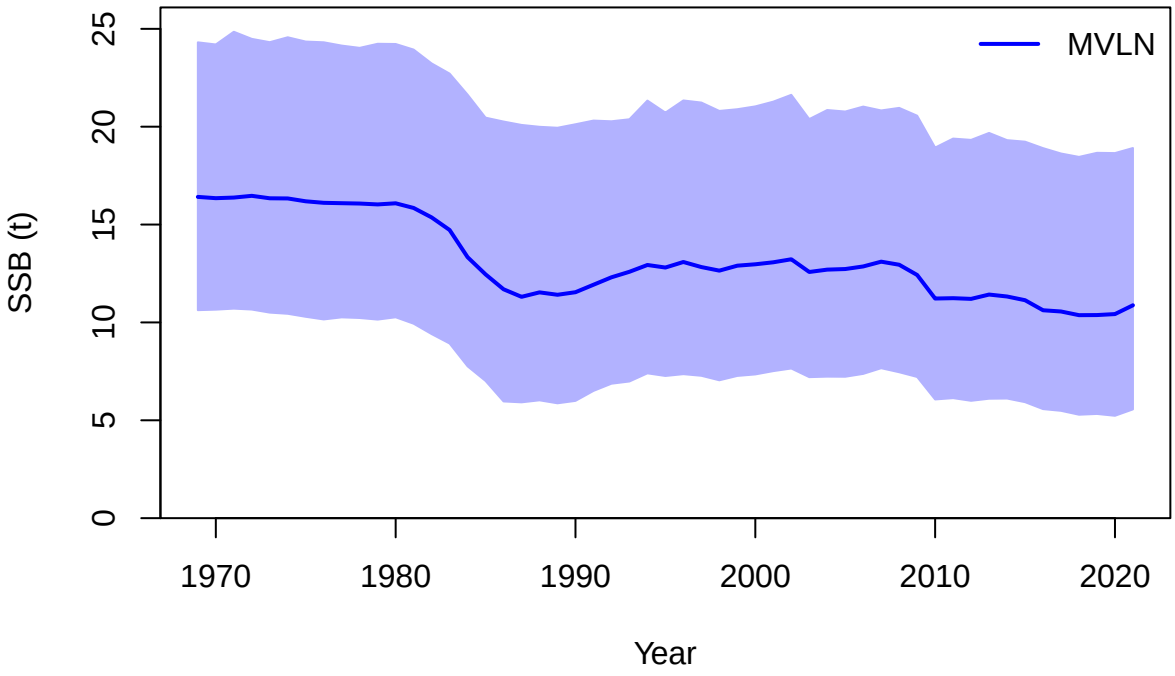

```
##
## Plot Comparison of stock
```



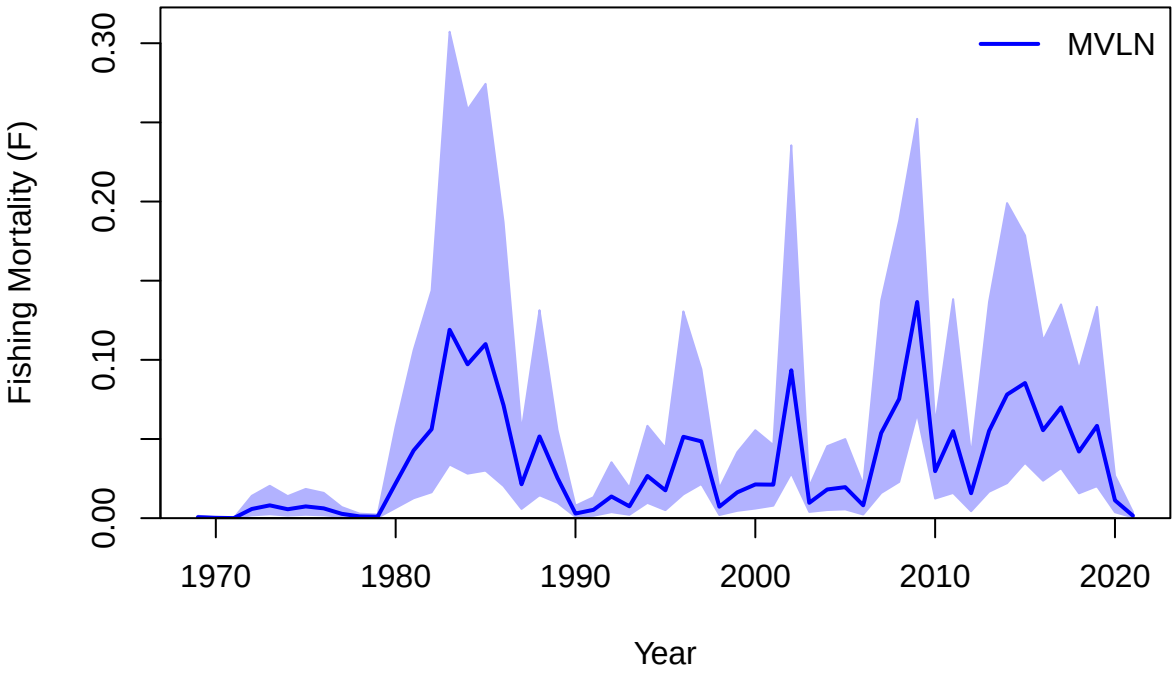

```
##
## Plot Comparison of harvest
```




```
##
## Plot Comparison of SSB
```

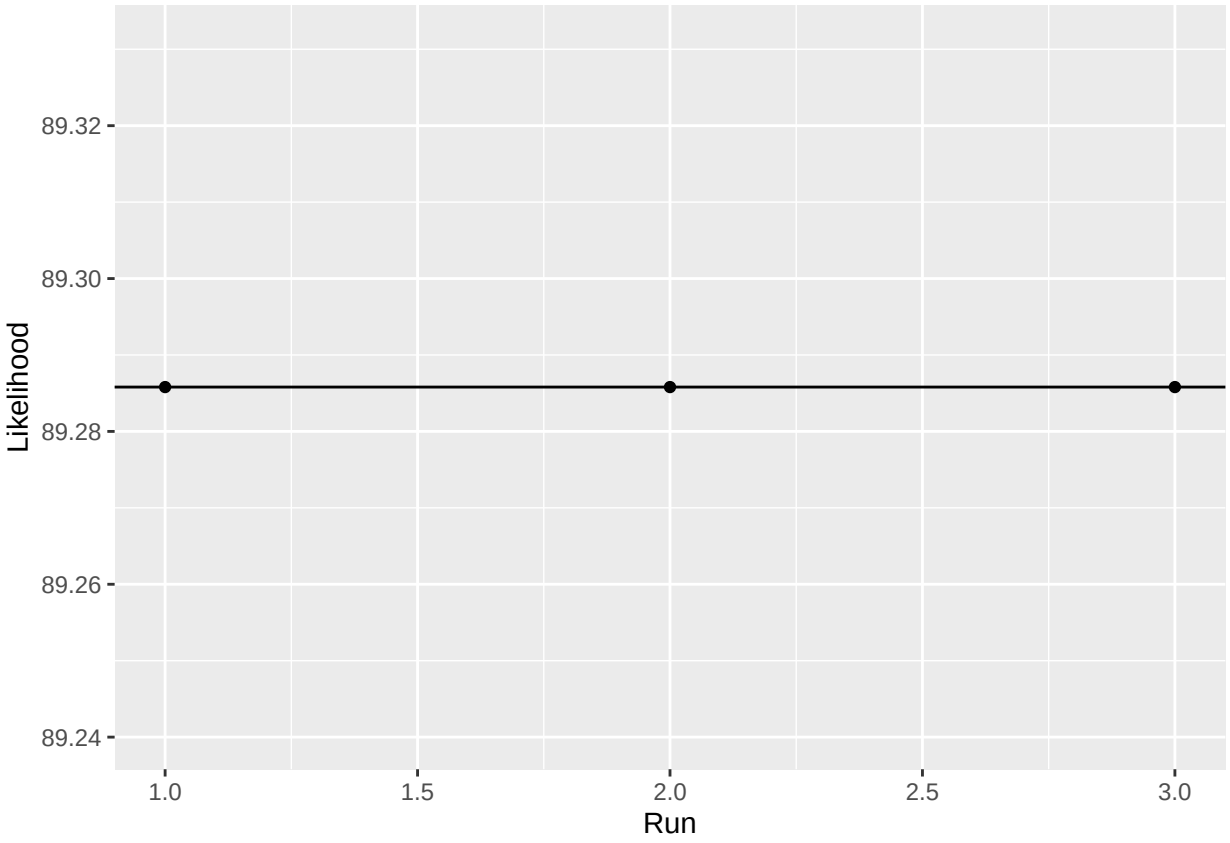


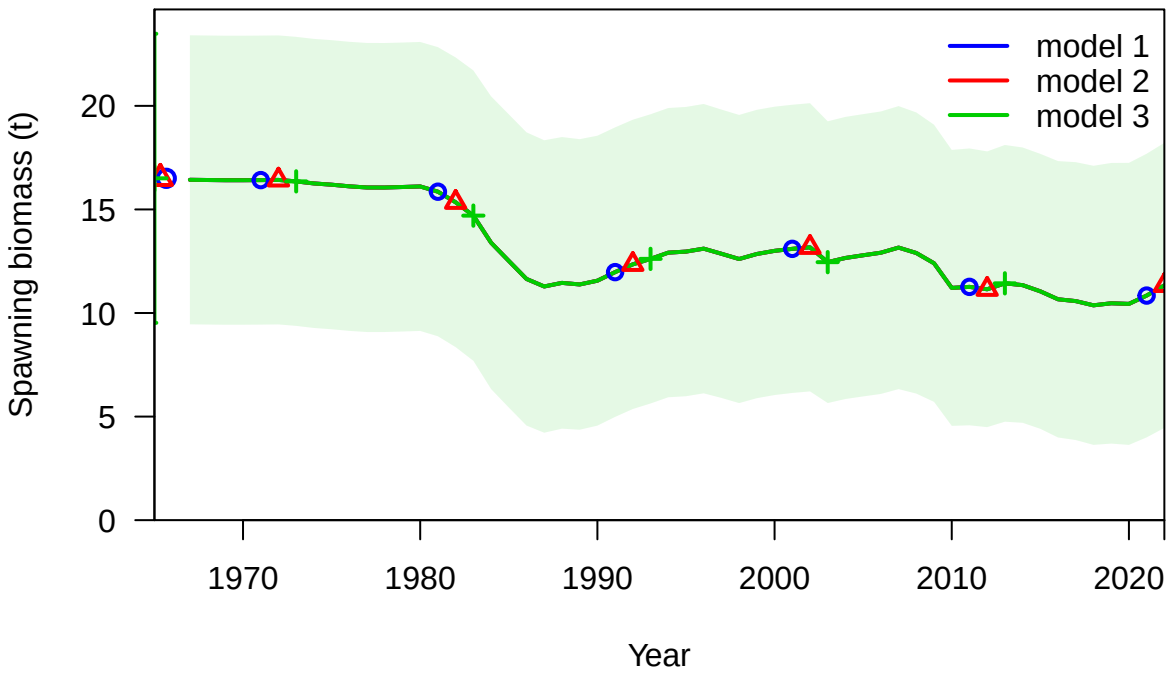

```
##
## Plot Comparison of F
```

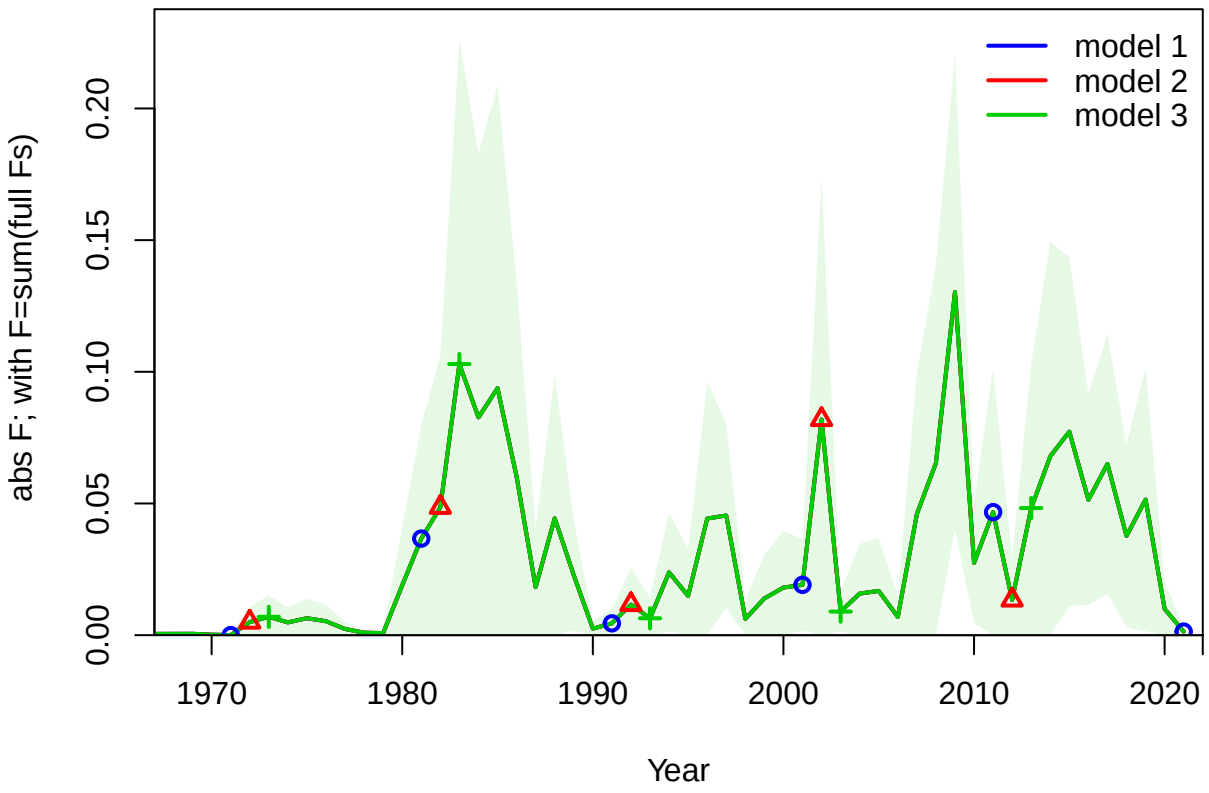

MVLN
Fishing Mortality (F)
0.00
0.10
0.20
0.30
1970
1980
1990
2000
2010
2020
Year

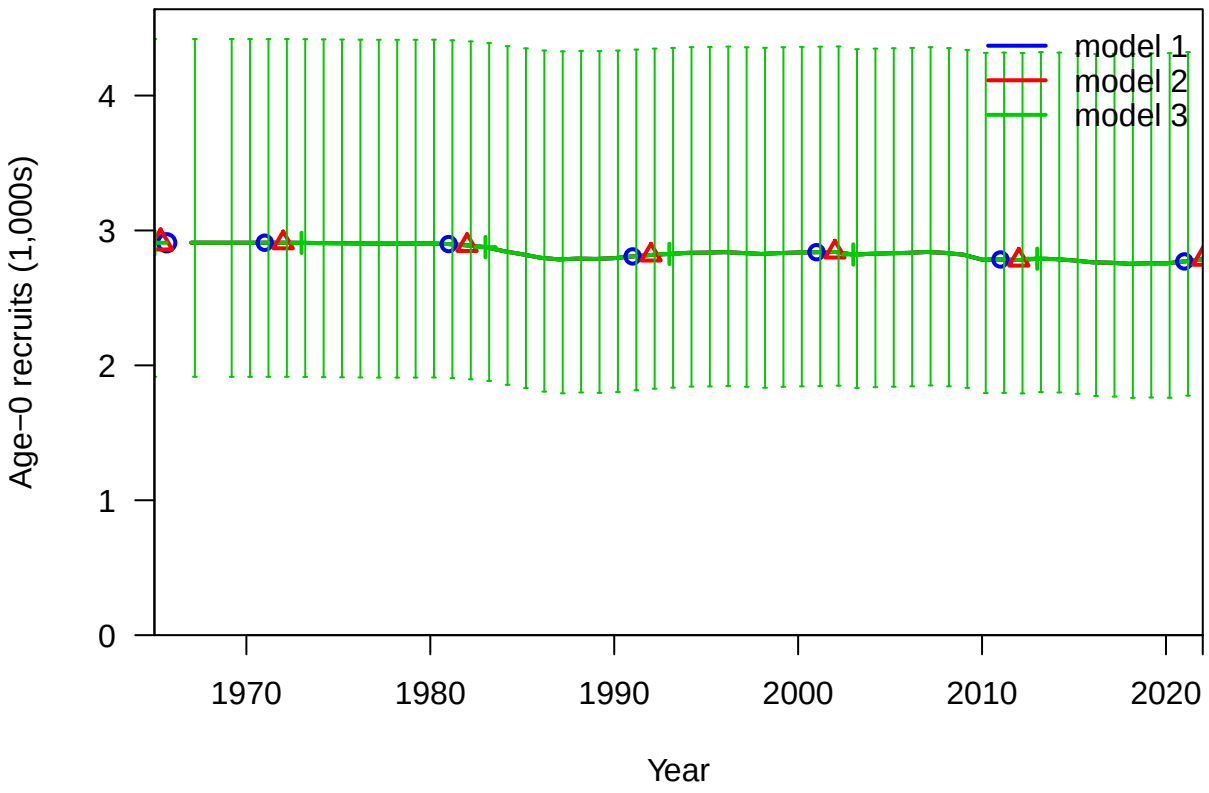
```
## RStudioGD
##         2
```

Jitter




model 1
model 2
model 3
Spawning biomass (t)
0
5
10
15
20
1970
1980
1990
2000
2010
2020
Year


model 1
model 2
model 3
abs F; with F=sum(full Fs)
0.00
0.05
0.10
0.15
0.20
1970
1980
1990
2000
2010
2020
Year


model 1
model 2
model 3
Age−0 recruits (1,000s)
0
1
2
3
4
1970
1980
1990
2000
2010
2020
Year