Dissect SAM: Catch observations

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SAM assumes

$$\begin{split} \log N_{1,y} &= \log R(\mathbf{N}_{y-1}) + \eta_{1,y} \\ \log N_{a,y} &= \log N_{a-1,y-1} - F_{a-1,y-1} - M_{a-1,y-1} + \eta_{a,y} \\ \log N_{A,y} &= \log (N_{A-1,y-1} e^{-F_{A-1,y-1} - M_{A-1,y-1}} + N_{A,y-1} e^{-F_{A,y-1} - M_{A,y-1}}) + \eta_{A,y} \end{split}$$

were

$$\log \mathbf{F}_y = \log \mathbf{F}_{y-1} + \boldsymbol{\xi}_y$$

Observe:

$$\log C_{a,y} = \log \left(\frac{F_{a,y}}{F_{a,y} + M_{a,y}} (1 - e^{-F_{a,y} - M_{a,y}}) N_{a,y} \right) + \epsilon_{a,y}^{c}$$

$$\log I_{y}^{(s)} = \log (Q_{a}^{(s)} e^{-(F_{a,y} + M_{a,y})day^{(s)}/365} N_{a,y}) + \epsilon_{a,y}^{s}$$

Assumes η_y , ξ_y and ϵ_y^C and ϵ_y^s all Gaussian distributed.



Observations

 Observations provides information about the system trough the observation equations:

$$\log C_{a,y} = \log \left(\frac{F_{a,y}}{F_{a,y} + M_{a,y}} (1 - e^{-F_{a,y} - M_{a,y}}) N_{a,y} \right) + \epsilon_{a,y}^{c}$$

$$\log I_{y}^{(s)} = \log \left(Q_{a}^{(s)} e^{-(F_{a,y} + M_{a,y}) day^{(s)}/365} N_{a,y} \right) + \epsilon_{a,y}^{s}$$

 When we have set up the system for F and N, we indirectly gain insight about those quantities trough observations and the observation equations.

Catch equation

We observe catch:

$$\log C_{a,y} = \log \left(\frac{F_{a,y}}{F_{a,y} + M_{a,y}} (1 - e^{-F_{a,y} - M_{a,y}}) N_{a,y} \right) + \epsilon_{a,y}^{c}$$

where $\epsilon_y^c \sim N(0, \Sigma_C)$.

Note

- $\frac{F_{a,y}}{F_{a,y}+M_{a,y}}$ is the proportion died in fishery
- $(1 e^{-F_{a,y}-M_{a,y}})N_{a,y}$ is the total amount of fish died

Exercise: Catch observations

- We will now investigate the implementation of the catch equation.
- Data about F and N are provided in Cobs.RData.
 - It may sound artificial that we know these quantities, but now we will now only focus on the catch equation.
- Exercise a) Implement the model

$$\log \textit{C}_{\textit{a},\textit{y}} \sim \textit{N}(\log \hat{\textit{C}}_{\textit{a},\textit{y}}, \sigma^2)$$

where

$$\begin{split} \log \hat{C}_{a,y} &= \log \left(\frac{F_{a,y}}{F_{a,y} + M_{a,y}} (1 - e^{-F_{a,y} - M_{a,y}}) N_{a,y} \right) \\ &= \log F_{a,y} - \log (F_{a,y} + M_{a,y}) + \log (1 - e^{-F_{a,y} - M_{a,y}}) + \log N_{a,y} \end{split}$$

• Exercise b) Include separate variance parameter per age.



Exercise: Catch observations

TMB output while fitting the model:

```
R
                                                                                              C++
> fit <- nlminb(obj$par, obj$fn, obj$gr)
outer mac:
             502.1106
            375.0361
outer mgc:
                                                       Data
outer mac:
             300.1951
                                                       Parameters
                                                                                          f(par,data) = ...
outer mac:
            16.76699
            8.446832
outer mgc:
outer mgc: 0.1371361
                                                       Optimization routine
                                                                                    Behind the scenes:
outer mgc:
            0.00109791
             1.442425e-07
outer mac:
>
```

Explain the output



Note

In the full model, optimal ${\bf F}$ and ${\bf N}$ are found in the inner optimization.

```
outer mgc: 2.125321
         158.72116: -0.405600 1.00234
iter: 1 value: 167.6788 mgc: 0.9840473 ustep: 1
iter: 2 value: 167.6786 mgc: 0.01624102 ustep: 1
iter: 3 value: 167.6786 mgc: 1.878016e-05 ustep: 1
iter: 4 mac: 3.701173e-11
                                                                                   R
iter: 1 value: 153.1634 mgc: 0.05411768 ustep: 1
iter: 2 value: 153.1634 mgc: 0.0001761833 ustep: 1
iter: 3 mgc: 1.925322e-09
                                                                           Data
iter: 1 value: 153.1634 mgc: 0.05411768 ustep: 1
                                                                           Parameters
                                                                                                            f(nar data) =
iter: 2 value: 153.1634 mgc: 0.0001761833 ustep: 1
iter: 3 mac: 1.925322e-09
outer mac: 0.4393079
                                                                           Optimization routine
                                                                                                      Rehind the scenes
         158.64804: -0.348985 1.00601
iter: 1 value: 150.4947 mgc: 0.2225958 ustep: 1
iter: 2 value: 150.4947 mgc: 0.0004934857 ustep: 1
iter: 3 mgc: 9.436551e-09
iter: 1 value: 152.9936 mgc: 0.006990456 ustep: 1
iter: 2 value: 152.9936 mgc: 6.445189e-07 ustep: 1
iter: 3 mac: 1.110223e-14
iter: 1 value: 152,9936 mgc: 0.006990456 ustep: 1
iter: 2 value: 152.9936 mgc: 6.445189e-07 ustep: 1
iter: 3 mac: 1.110223e-14
outer mgc: 0.01172901
```

Catch observation configurations in SAM

- ullet Use keyVarObs to modify the catch observation variance ($\Sigma_{a,a}$)
- First line is typically the catch
- Example with same variance parameter for all ages:

Catch observation configurations in SAM

 We can construct a link between predicted observations and associated variance in SAM

Let $\mu_{a,y}$ be the predicted observation for age a at year y on natural scale, and let $v_{a,y}$ be the corresponding variance. With this option we impose the assumption that

$$\mathbf{v}_{\mathbf{a},\mathbf{y}} = \alpha \mu_{\mathbf{a},\mathbf{y}}^{\beta},$$

and estimate α and β internally in SAM.

- ullet keyVarObs couples the lpha parameters.
- ullet predVarObsLink couples the eta parameters.

Exercise: Include this structure in the previous exercise

• Hint: Variance of log observation is given by:

$$\sigma_{a,y}^2 = \log\left(\alpha_a \mu_{a,y}^{\beta_a - 2} + 1\right).$$



External observation variances in SAM

- ullet We may know an external estimate of the uncertainty of $\log \hat{\mathcal{C}}_{a,y}$
- We can include such data as a relative weighting factor for each observation
- See the vignette obsCovarOptions for details



Exercise: Include external covariance structures

A script for fitting SAM to North East Arctic cod is provided in the folder externalCovarianceEx.

- Externally estimated log catch covariance matrices are provided in covCatch.RData
- A SAM script without using external variances are provided in script.R
- Exercise a: Read the vignette obsCovarOptions and include the external covariance matrices in the assessment.
- Exercise b: Based on AIC, do we obtain an improvement?

Total catches

- We inform the system with observations trough observations equations.
- Need to express observations as realisations of a probability distribution with parameters provided by parameters in the model (e.g. F and N).
- Say we only observe total catch, and not per age.
- We can include that observation trough:

$$\log C_{y} = \log \left(\sum_{a} \frac{F_{a,y}}{F_{a,y} + M_{a,y}} (1 - e^{-F_{a,y} - M_{a,y}}) N_{a,y} cw_{a,y} \right) + \epsilon_{y}^{C}$$

where $\epsilon_{\it V}^{\it C} \sim N(0, \sigma_{\it V}^2)$.

• Here $cw_{a,y}$ is mean catch weight at age a in year y.



Total catches

• Can use the delta method for constructing variance of $\epsilon_{\it y}^{\it Catch}$:

$$\sigma_y^2 = h_y^t \Sigma_C h_y$$

where

$$h_{y} = (\frac{\hat{C}_{1,y}cw_{1,y}}{\sum_{a}\hat{C}_{a,y}cw_{a,y}}, \cdots, \frac{\hat{C}_{A,y}cw_{A,y}}{\sum_{a}\hat{C}_{a,y}cw_{a,y}})$$

and

$$\hat{C}_{A,y} = \frac{F_{a,y}}{F_{a,y} + M_{a,y}} (1 - e^{-F_{a,y} - M_{a,y}}) N_{a,y}.$$

- We can now include total catch in years when catch at age is not available.
- Procedure in SAM: Include total catch as a survey (of age -1), set
 keyBiomassTreat =3 and set catch per age to -1 in the corresponding years
- Similarly we can include total landings (keyBiomassTreat = 4)



Biomass indices

- Observations of total spawning stock biomass
- SSB observations can be represented internally as:

$$\log {\sf SSB}_y = \log \Big(\sum_{a} Q_a^{(s)} e^{-(F_{a,y} + M_{a,y}) day^{(s)}/365} N_{a,y} m_{a,y} w_{a,y} \Big) + \epsilon_y^{SSB}$$

where $m_{a,y}$ is proportion mature and $w_{a,y}$ is mean stock weight.

- Procedure in SAM: Including as survey of age -1 and set keyBiomassTreat = 0.
- Similarity we can include:
 - Catch index (keyBiomassTreat = 1)
 - FSB index (keyBiomassTreat = 2)
 - TSB index (keyBiomassTreat = 5)

