

# Status Report: Gmacs BBRKC

Darcy Webber • Jim Ianelli  
January 2017

# Gmacs SMBKC

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A generic size-structured stock assessment model <http://seacode.github.io/gmacs>

1,322 commits 3 branches 4 releases 6 contributors

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| Author     | Commit Message    | Time                                                               |
|------------|-------------------|--------------------------------------------------------------------|
| quantifish | delete twosex dir | Latest commit 3e4bc7 19 minutes ago                                |
|            | docs              | delete twosex dir 18 minutes ago                                   |
|            | examples          | delete twosex dir 18 minutes ago                                   |
|            | gnr               | delete twosex dir 18 minutes ago                                   |
|            | src               | comparsions all set up 8 hours ago                                 |
|            | .gitignore        | backed off weight on NMFS trawl survey 6 days ago                  |
|            | .travis.yml       | Minor edit to travis.yml for email notifications 2 years ago       |
|            | CHANGELOG.md      | took a little bit to get the onesex model to fit again 2 years ago |
|            | Makefile          | tidy some old files 3 months ago                                   |
|            | README.md         | updates to readme and smbkc document 5 months ago                  |

## README.md

### Gmacs

A generalized size-structured stock assessment modelling framework. Gmacs includes:

- The Gmacs model
- A simulation model
- An R package for working with Gmacs output files
- A Wiki

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Releases Tags Draft a new release

**SMBKC 2016** Edit

Pre-release

quantifish released this on 28 Sep 2016 · 47 commits to develop since this release

This tag identifies the code that was used for the 2016 SMBKC stock assessment. This is the assessment after it was presented to the CPT and some edits were made to the document, but before the assessment was presented to the SSC.

### Downloads

- Source code (zip)
- Source code (tar.gz)

# Gmacs BBRKC Progress

- Sex-specific recruitment - uses a proportion m/f parameter that is logit transformed with normal prior (rather than 50/50 split)
- Sex-specific time-varying natural mortality
- Sex-specific custom growth matrices
- Sex-specific natural mortality rates by year (can be fixed-custom)
  
- Better numbers at length plots, selectivity plots, molt probability plots, etc
  
- Updated BBRKC model input data
- BBRKC model progress
- BBRKC document with comparisons in progress

Bristol Bay Red King Crab Stock Assessment 2017

D'Arcy Webber<sup>1</sup>, Jie Zheng<sup>2</sup>, and James Ianelli<sup>3</sup>

<sup>1</sup>Quantifish, darcy@quantifish.co.nz

<sup>2</sup>Alaska Department of Fish and Game, jie.zheng@alaska.gov

<sup>3</sup>NOAA, jim.ianelli@noaa.gov

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## Executive Summary

1. **Stock:** Red king crab (RKC), *Paralithodes camtschaticus*, in Bristol Bay, Alaska.

2. **Catches:** Peak historical harvest was 4288 tonnes (9.454 million pounds) in 1983/84<sup>1</sup>. The fishery was closed for 10 years after the stock was declared overfished in 1999. Fishing resumed in 2009/10 with a fishery-reported retained catch of 209 tonnes (0.461 million pounds), less than half the 529.3 tonne (1.167 million pound) TAC. Following three more years of modest harvests supported by a fishery catch per unit effort (CPUE) of around 10 crab per pot lift, the fishery was again closed in 2013/14

# Problems Encountered

## Gmacs

- Gmacs is slow (~15 minutes per BBRKC model run) - model flexibility adds excess baggage, be careful what you wish for
- Positive definite Hessian issues

## 2016 Model

- The 2016 model is initialised with no oldshell male crab in 1975. In 1976 they appear!
- The code is very difficult to follow. Figures in document can be misleading.

## Gmacs BBRKC

- Had to fix initial numbers at those estimated in 2016 model for now as I cannot match this initialisation - this likely causes other problems
- Had to fix growth matrix to 2016 model matrix for now as I could not match - 2016 model derived differently
- Poor fit to NMFS survey - particularly the bulge in biomass around 1990
- Not fitting to BSFRF survey if  $q = 1$

# Model Dimensions & Why Gmacs is Slow

|                  |    |                    |
|------------------|----|--------------------|
| Size-classes     | 20 | 65-165             |
| Sexes            | 2  | Male, Female       |
| Shell conditions | 2  | Oldshell, Newshell |
| Seasons          | 4  | 1-4                |
| Years            | 42 | 1975-2016          |

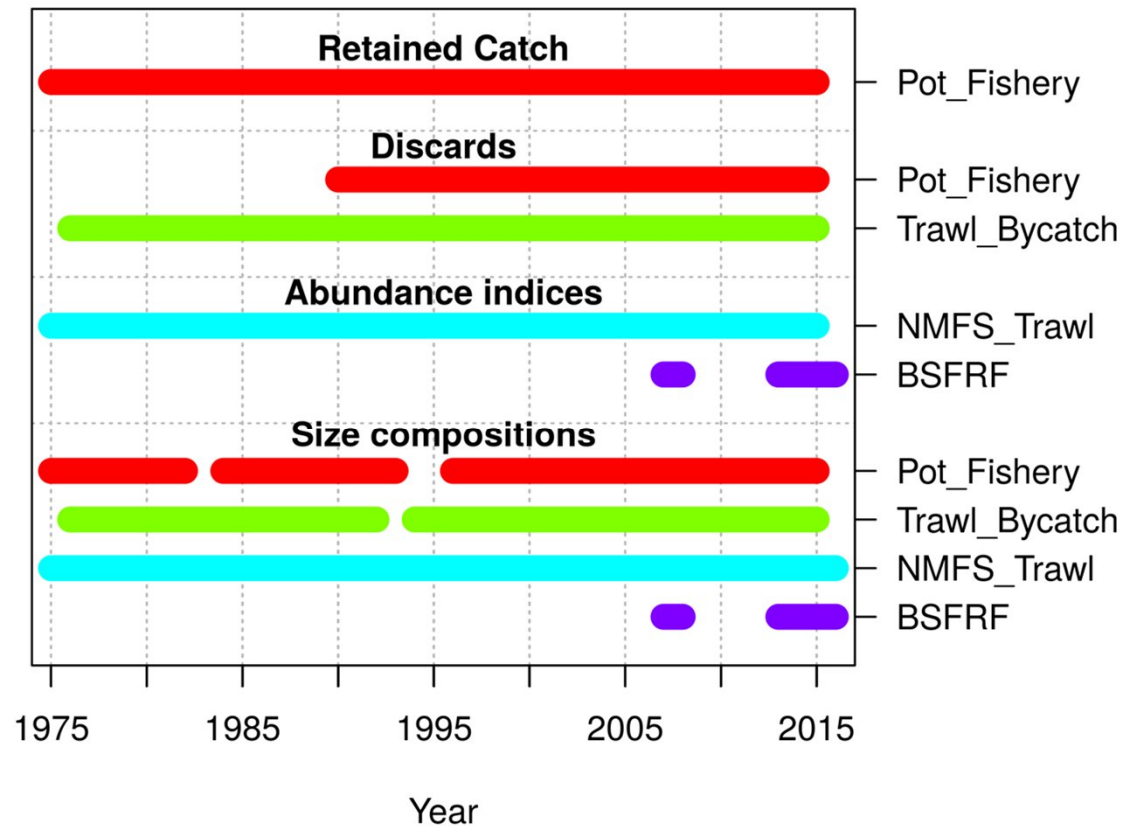
$20 \times 2 \times 2 \times 4 \times 42 = 13440$  dimensions

c.f. a BBRKC-specific model

$20 \times 3 \times 1 \times 42 = 2520$  dimensions

# Data

1. Catch
  - a. Pot fishery retained males
  - b. Pot fishery discarded males
  - c. Pot fishery discarded females
  - d. Trawl bycatch males+females
2. Survey
  - a. NMFS survey males
  - b. NMFS survey females
  - c. BSFRF survey males+females
3. Length-frequency
  - a. Pot fishery retained males
  - b. Pot fishery discarded males
  - c. Pot fishery discarded females
  - d. Trawl bycatch males
  - e. Trawl bycatch females
  - f. NMFS survey males
  - g. NMFS survey females
  - h. BSFRF survey males
  - i. BSFRF survey females



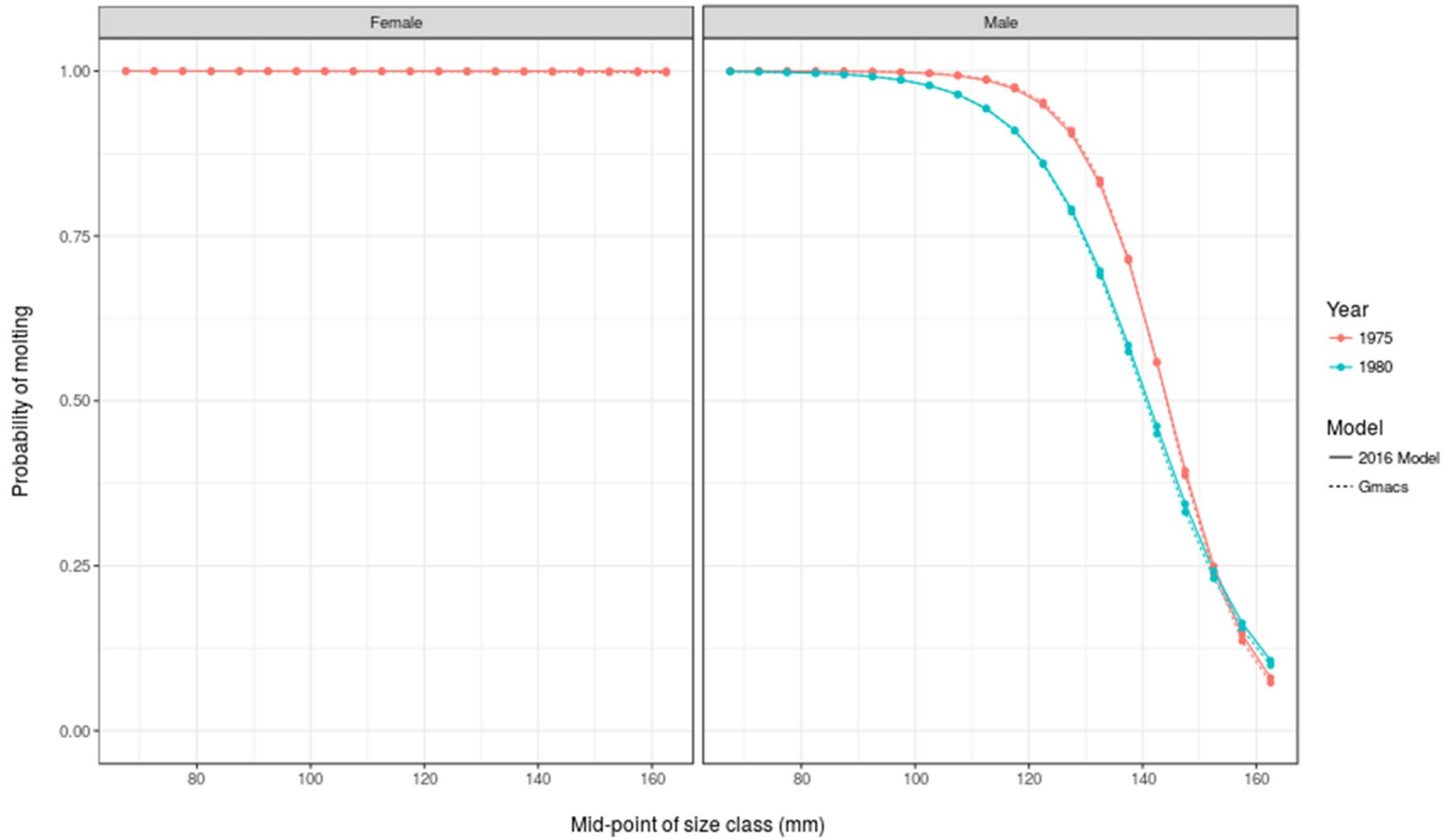
# Timing

Four seasons defined to try to best match 2016 model dynamics

| Season | What happens                                                                        |
|--------|-------------------------------------------------------------------------------------|
| 1      | Recruitment                                                                         |
| 2      | Trawl bycatch fishery                                                               |
| 3      | Natural mortality, molting & growth, directed pot fishery, surveys (NMFS and BSFRF) |
| 4      | Calculate MMB                                                                       |

# Molt Probability

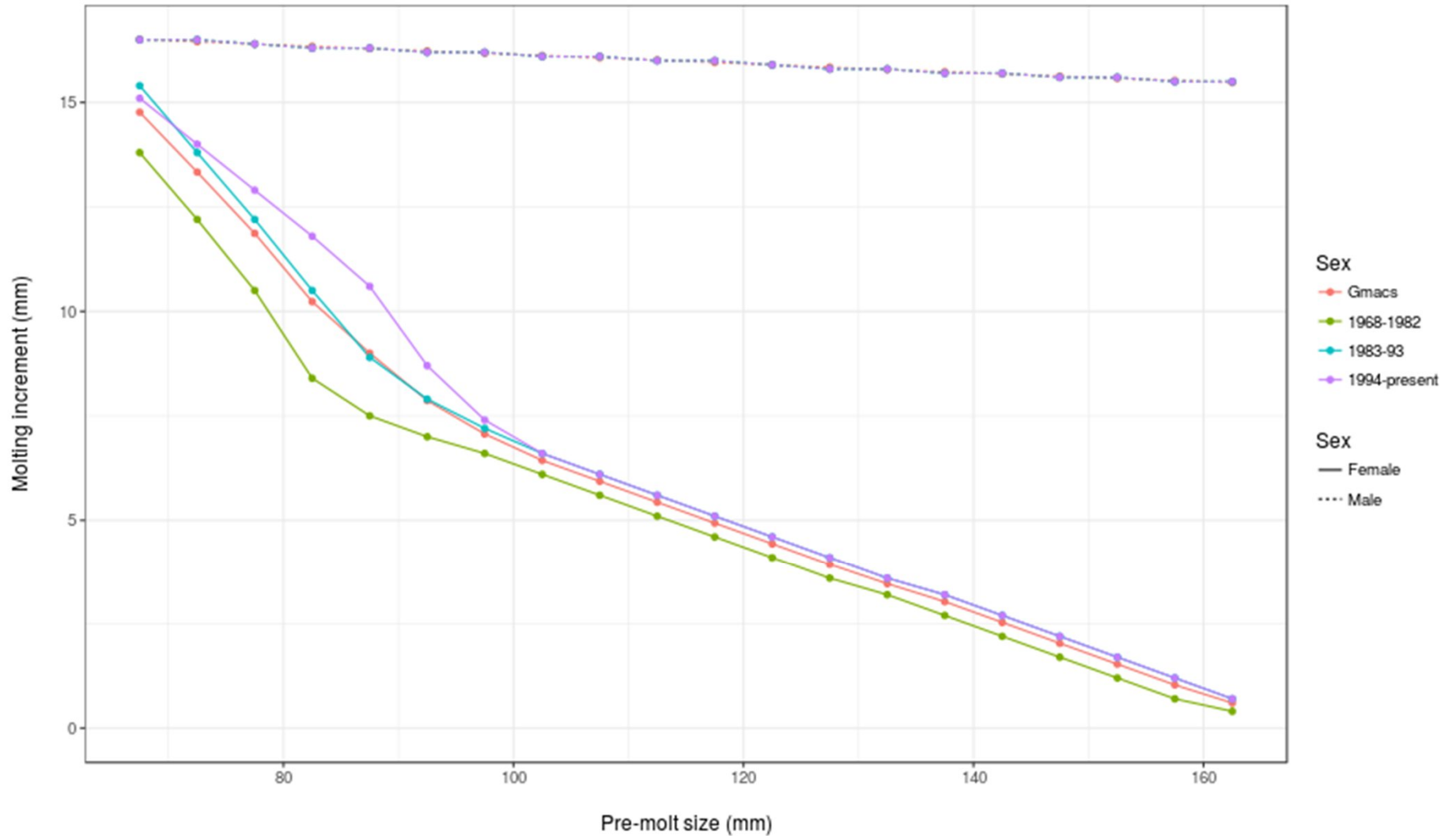
- Females molt every year
- Gmacs is using time-varying molt probability - these match up well with 2016 model





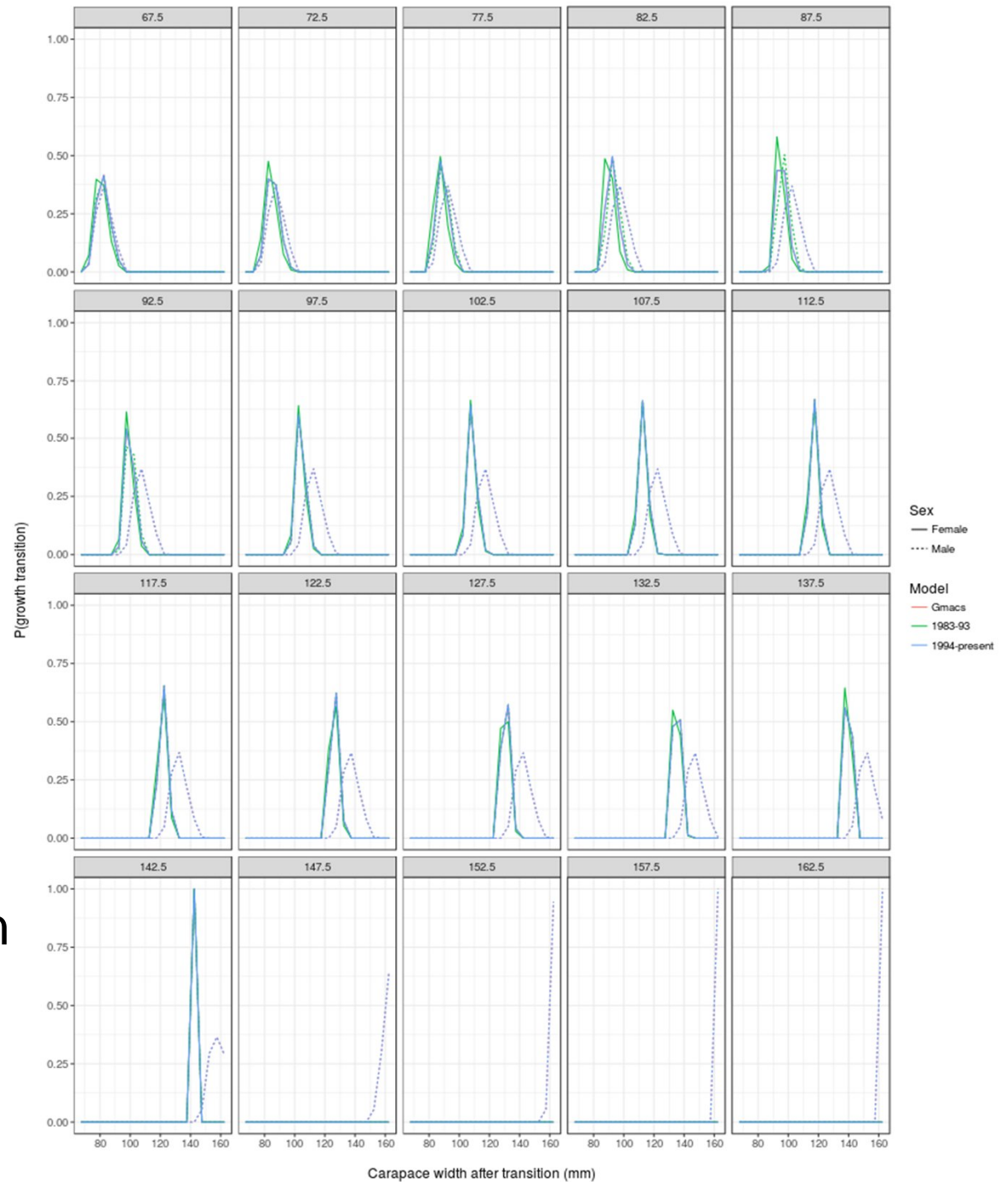
# Growth Increment Each Molt

- Gmacs not using time-varying growth (for females), as there is little evidence to support doing so

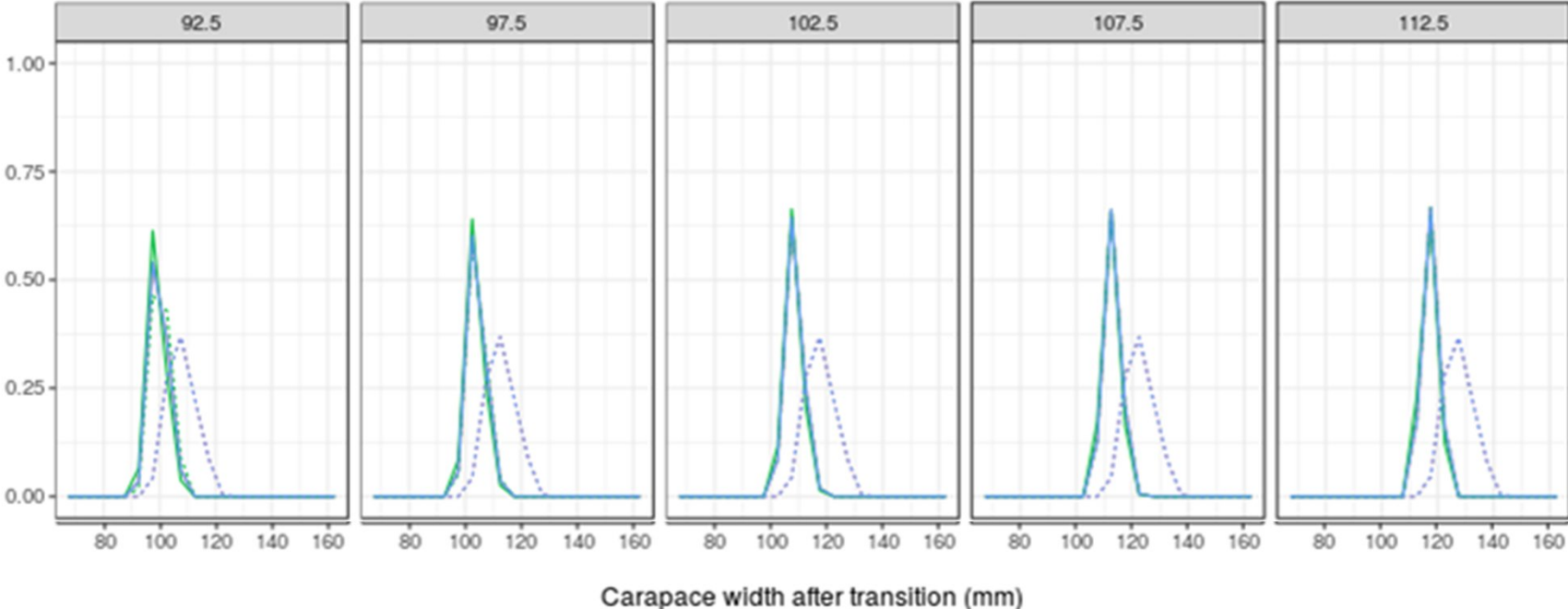


# Growth Matrix

- Gmacs not using time-varying growth (for females), as there is little evidence to support doing so
- Gmacs has fixed the growth matrix to the 2016 model as a similar matrix could not be derived (given the growth increments on the previous slide)

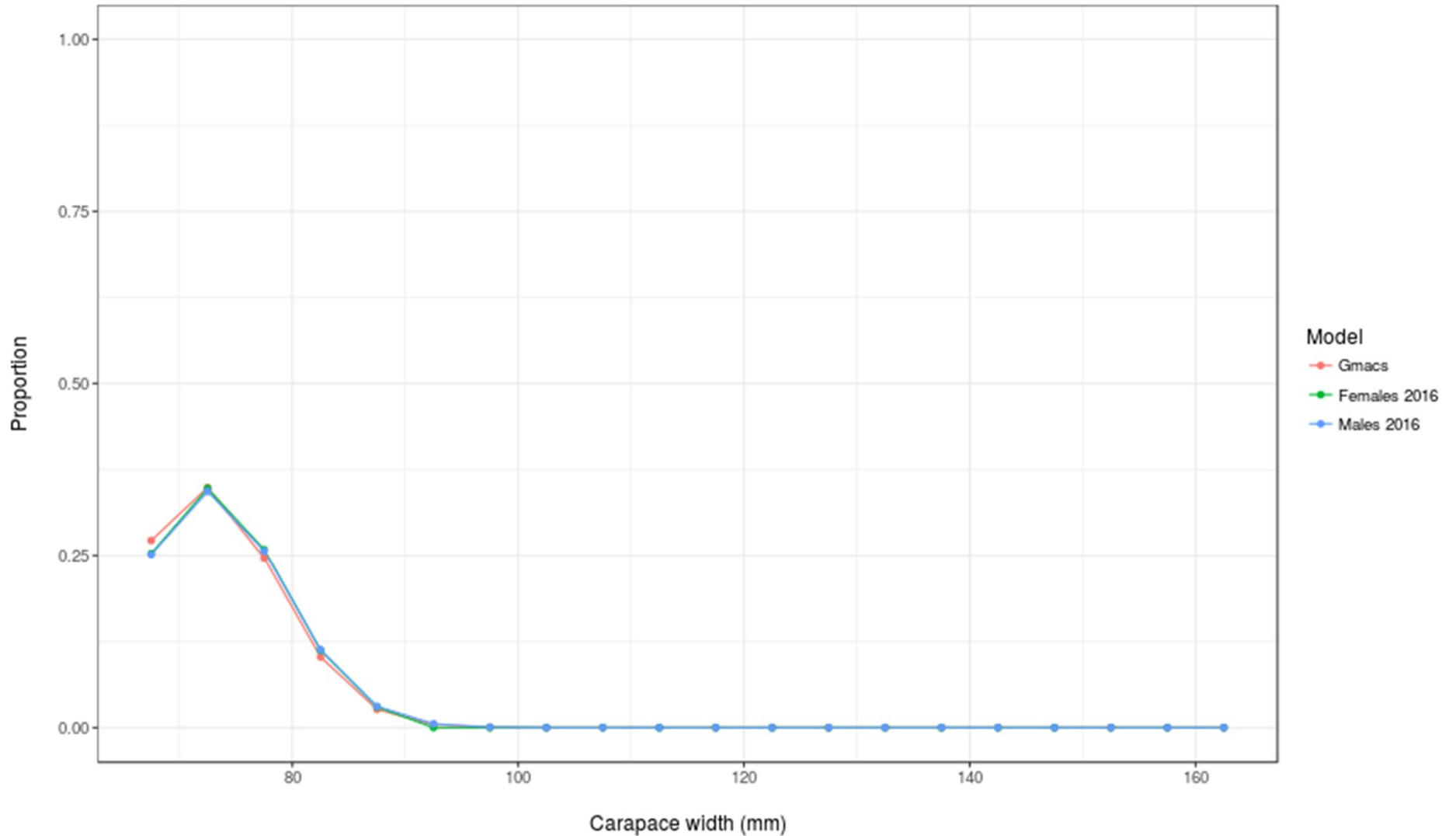


# Growth Matrix



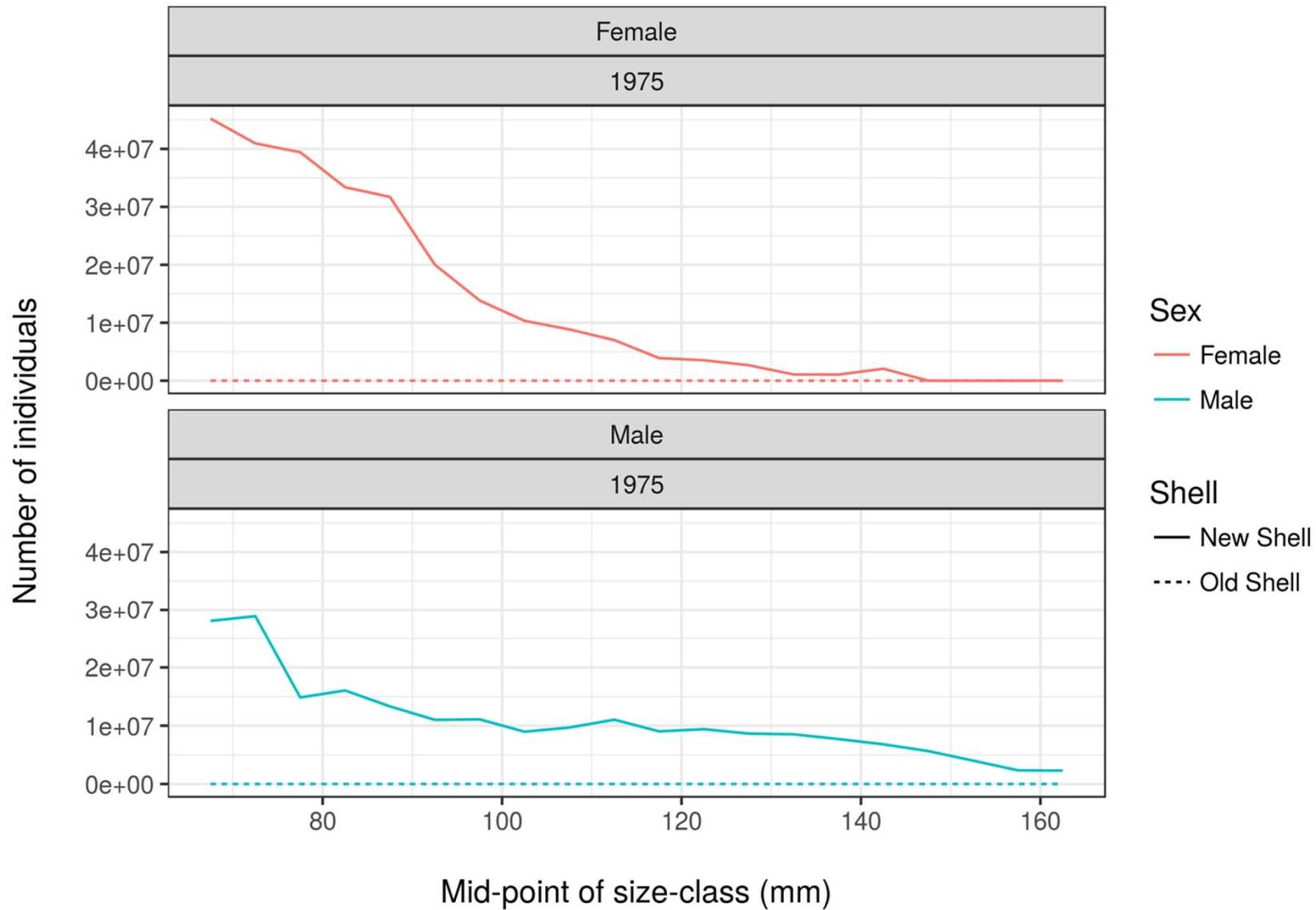
# Recruitment Size

Gmacs not using sex-specific recruitment size--seems unnecessary



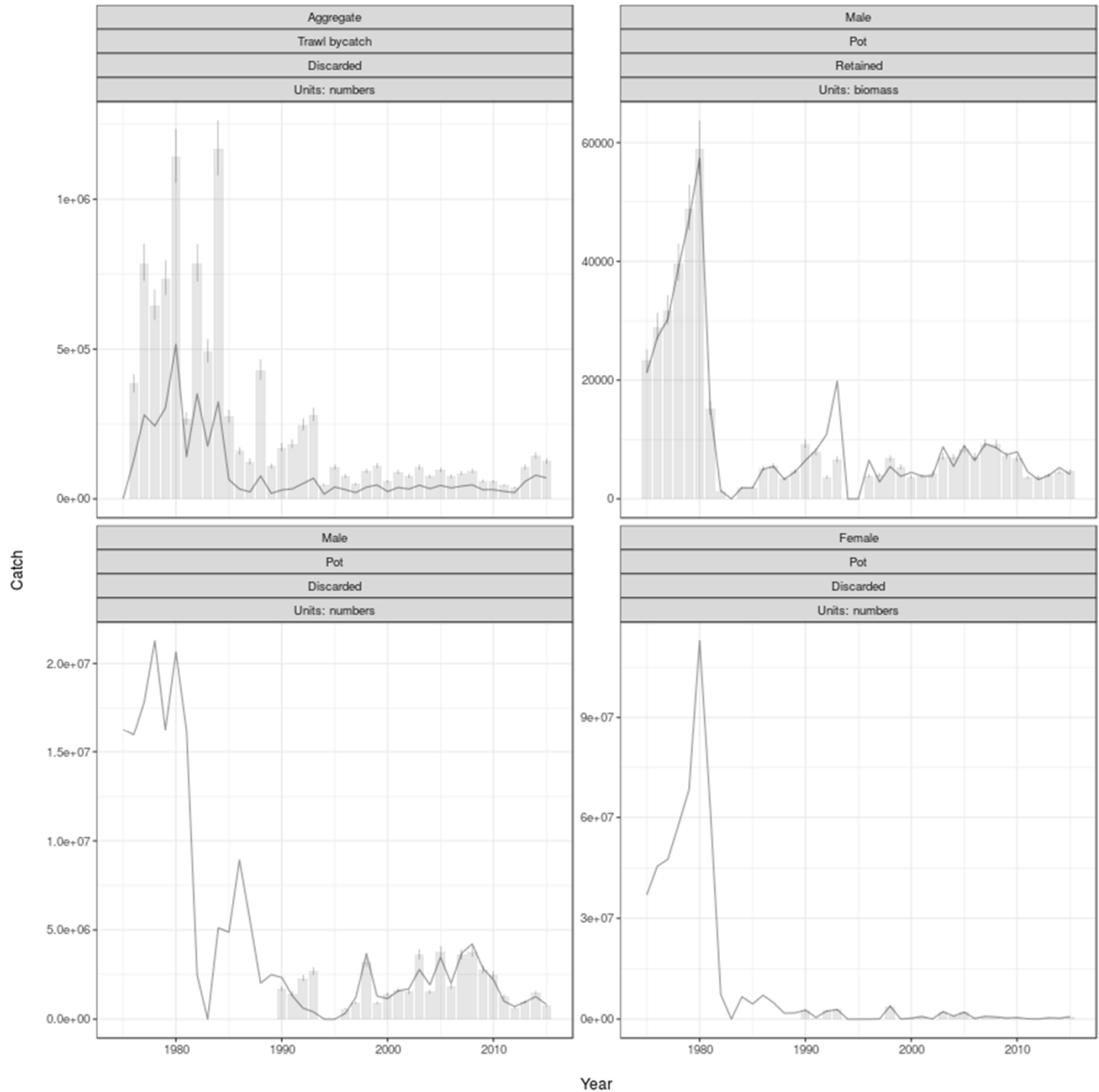
# Initial Numbers (season 1)

Gmacs initial numbers fixed at those used in 2016 model - could not replicate this initialisation - but this seems to be causing problems



# Catch

- Why is current model not using full catch time-series?



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# **Gmacs BBRKC: Different Model Structures**

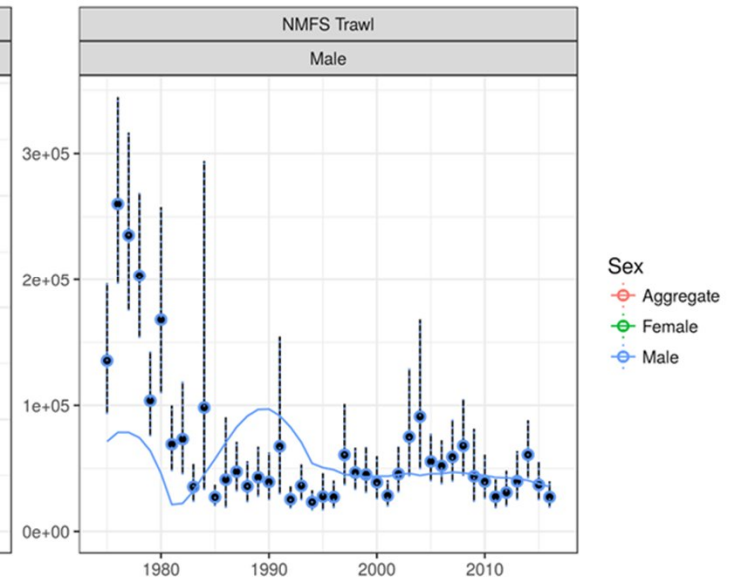
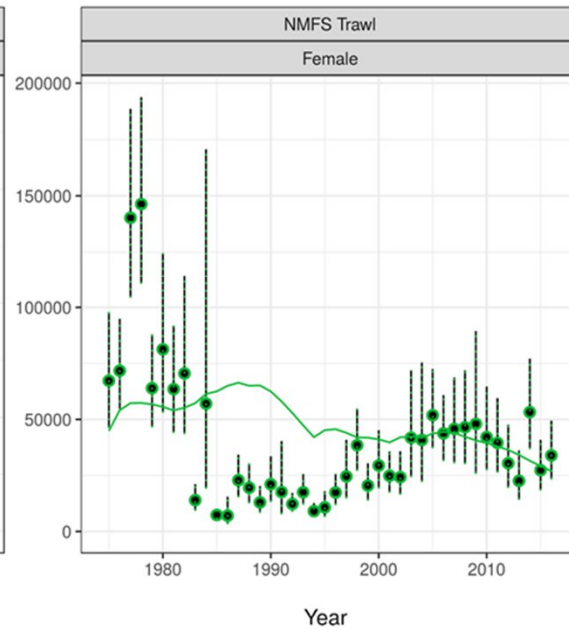
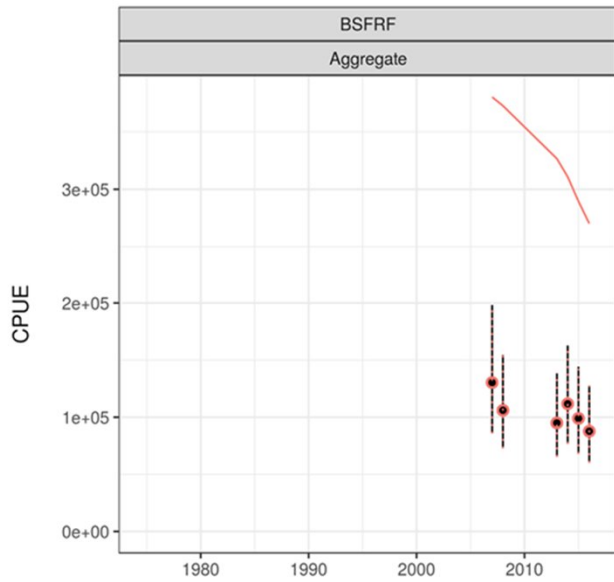
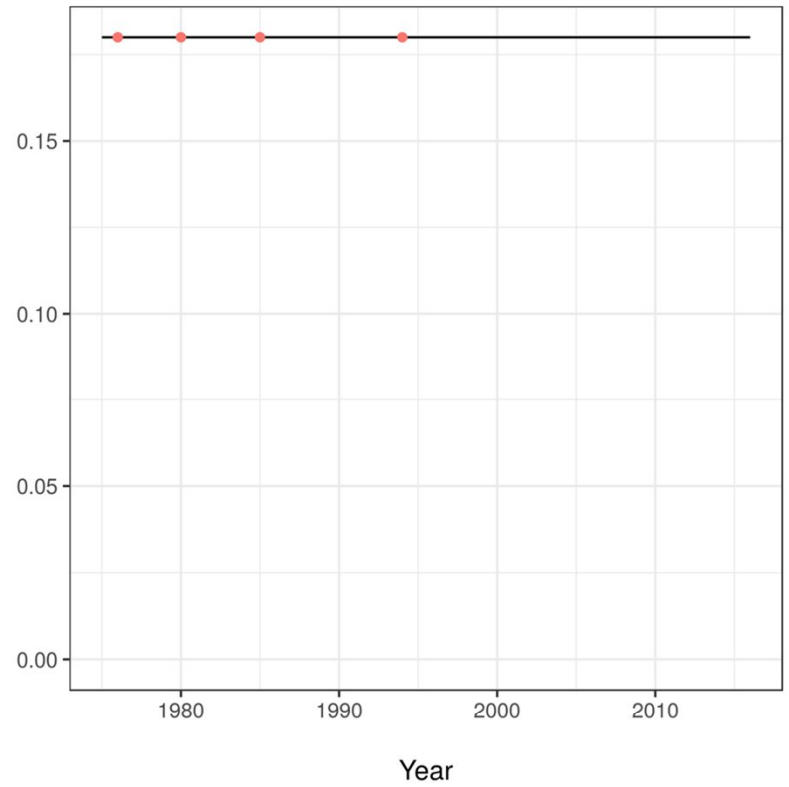
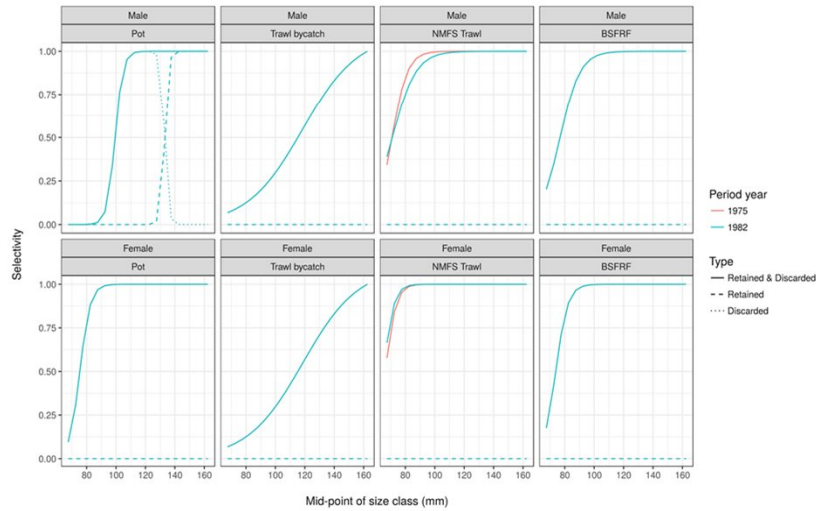
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# Different model structures

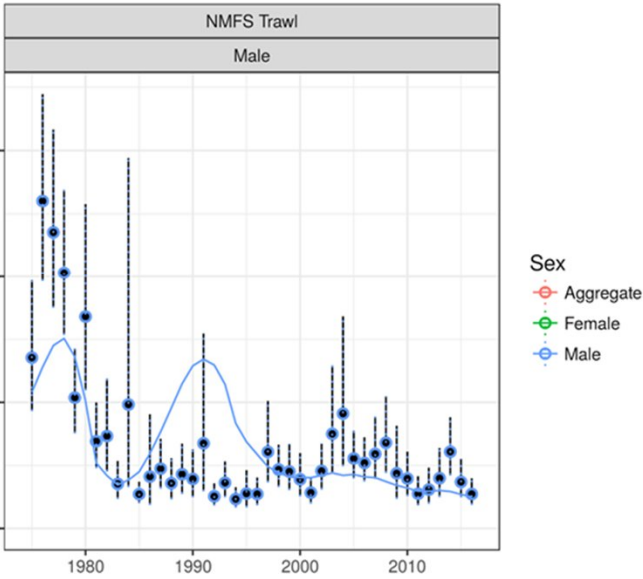
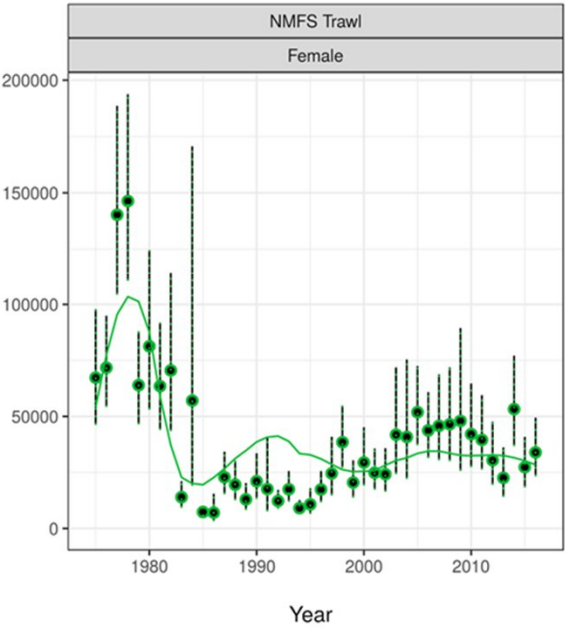
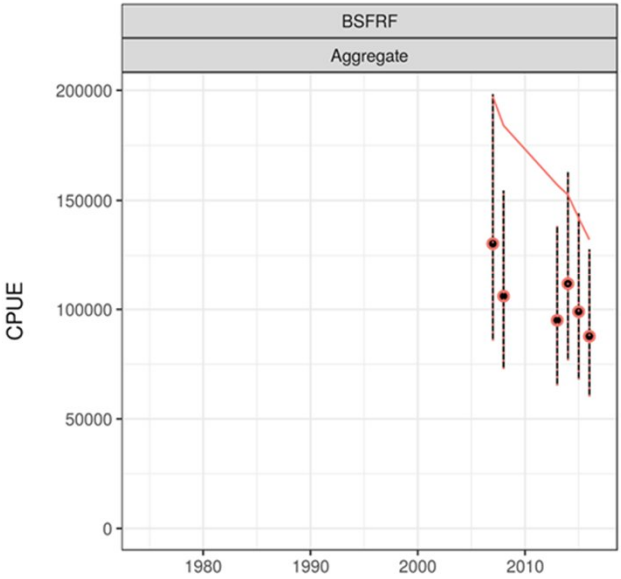
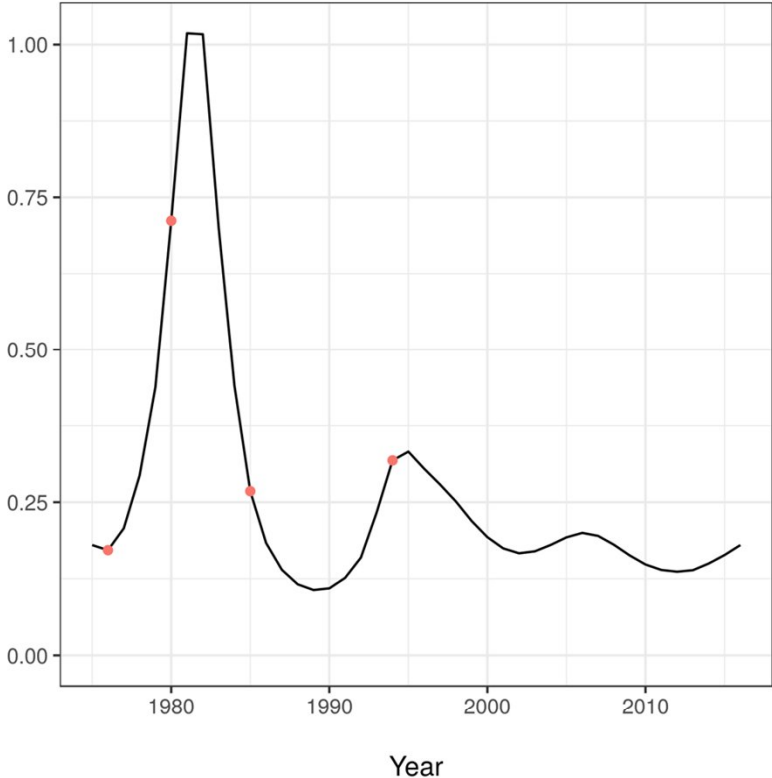
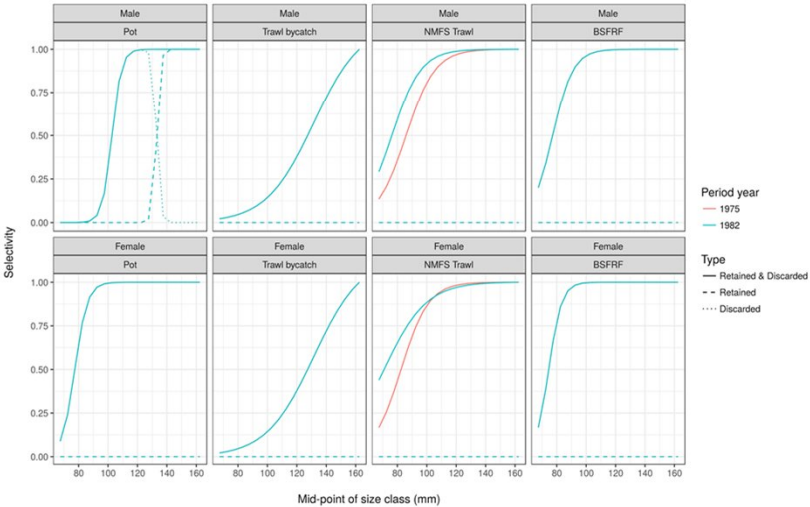
| <b>Model name</b> | <b>M</b>       | <b>BSFRF q</b> | <b>NMFS Lambda</b> |
|-------------------|----------------|----------------|--------------------|
| Constant M        | Constant       | Fixed at 1     | 1.0                |
| Random Walk M     | Random walk    | Fixed at 1     | 1.0                |
| Model M           | At 2016 values | Fixed at 1     | 1.0                |
| Estimate BSFRF q  | At 2016 values | Estimated      | 1.0                |
| NMFS Lambda=4     | At 2016 values | Fixed at 1     | 4.0                |



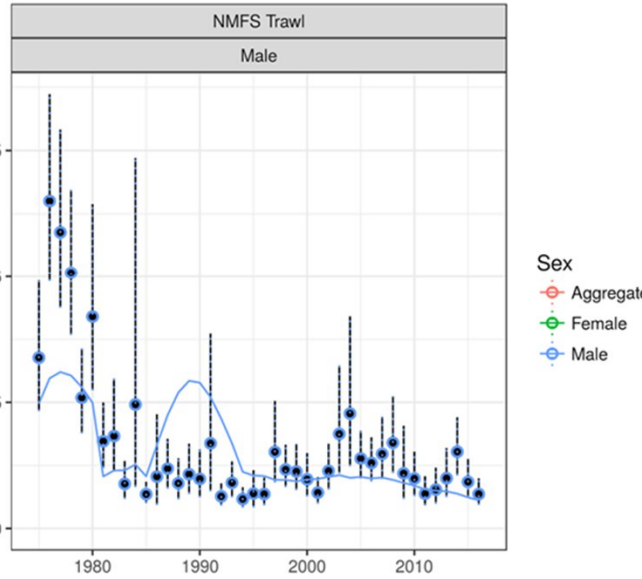
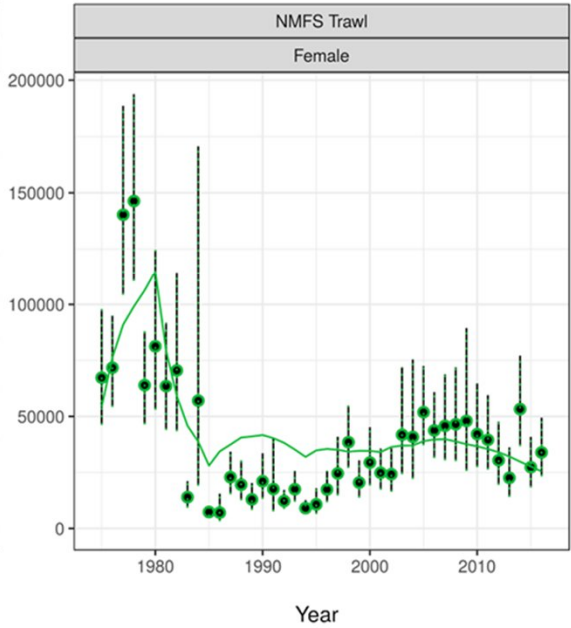
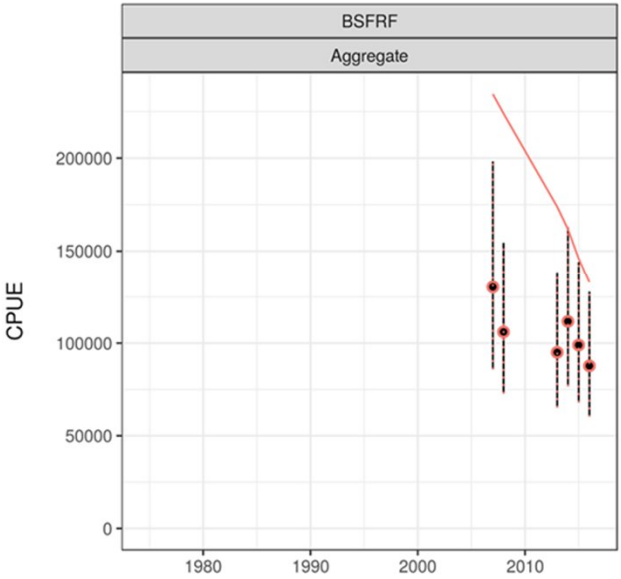
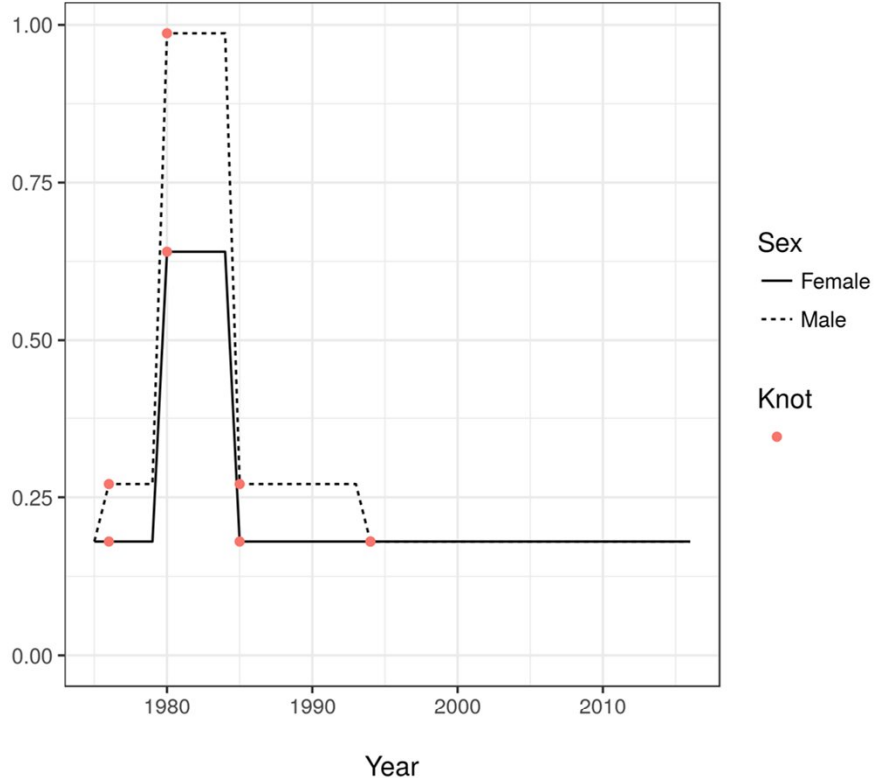
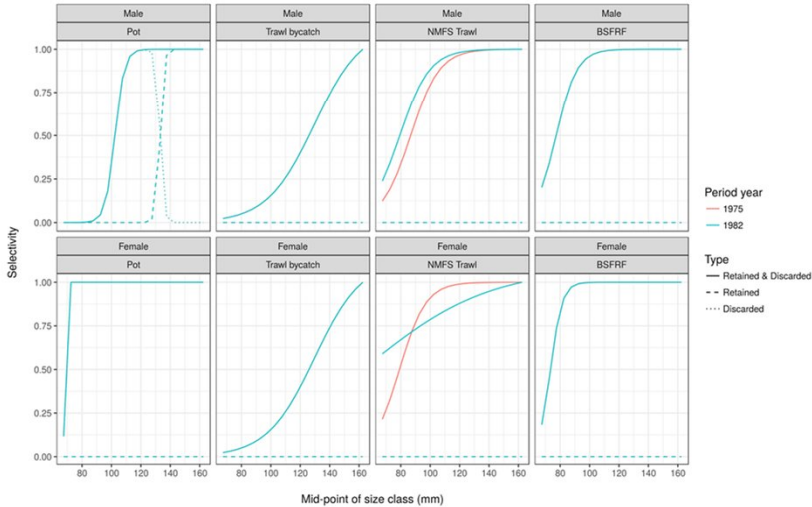
# 1.Constant M



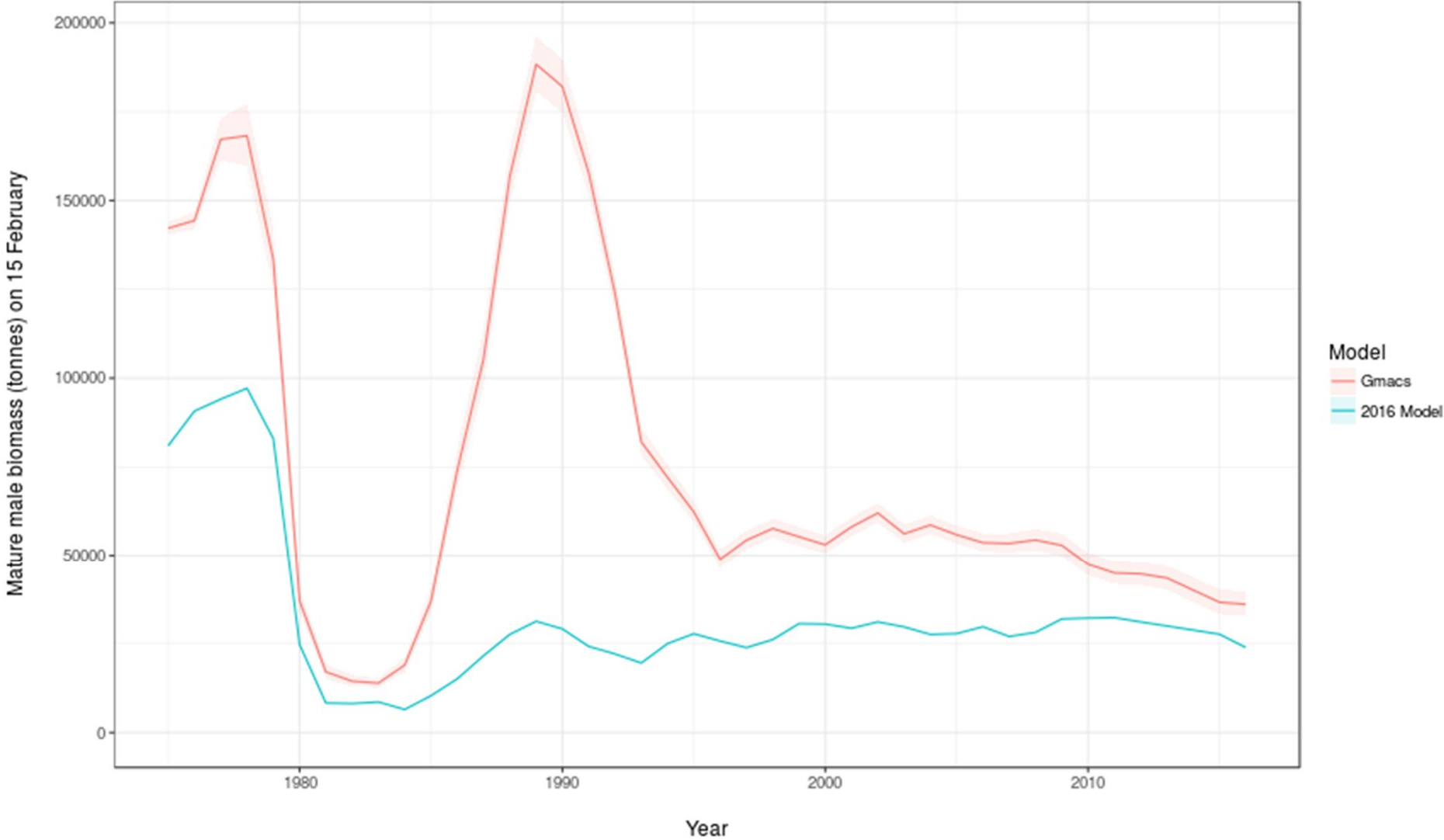
# 2. Random Walk M



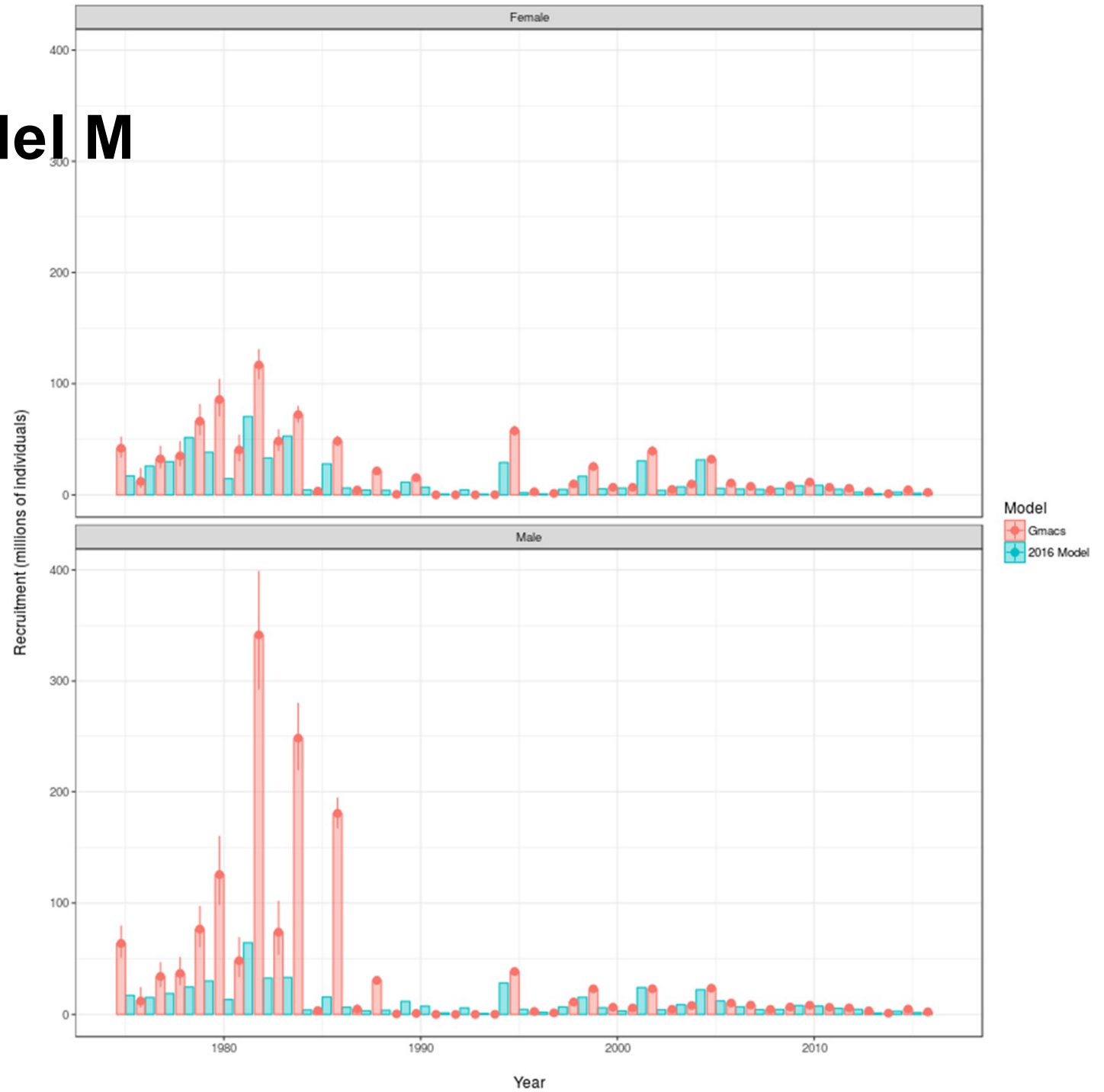
# 3. 2016 Model M



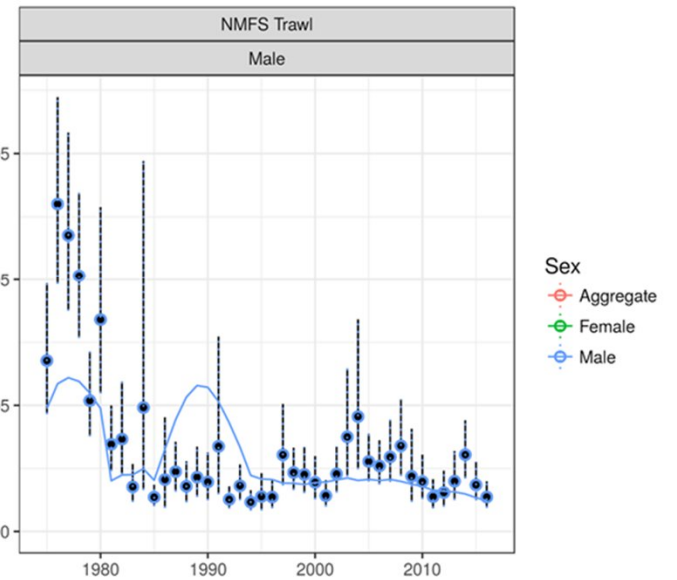
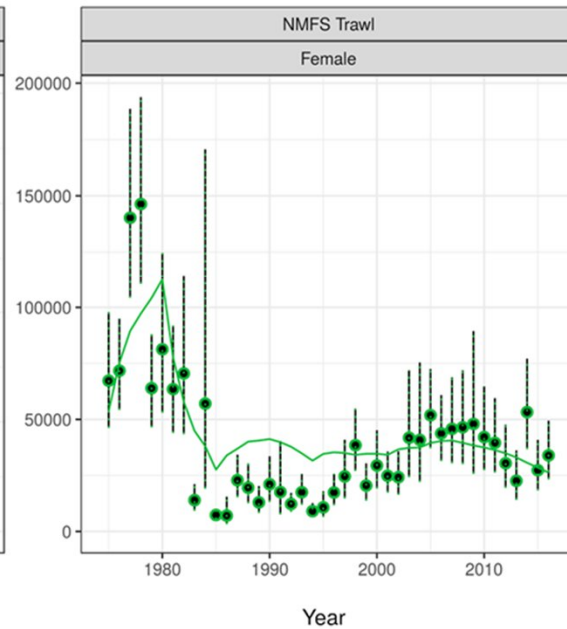
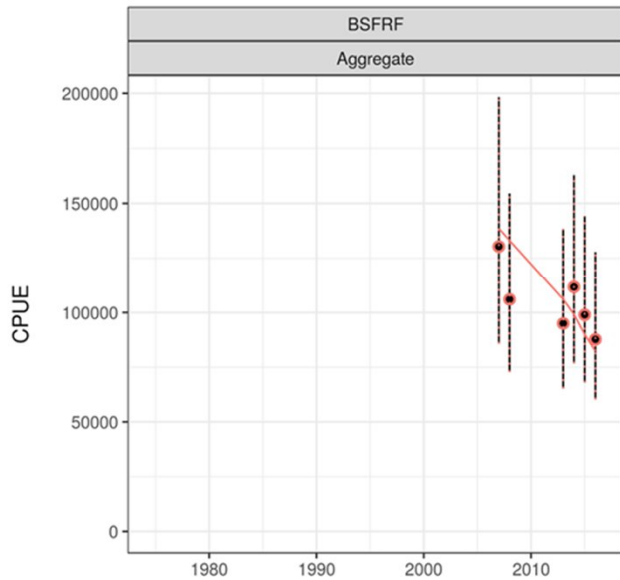
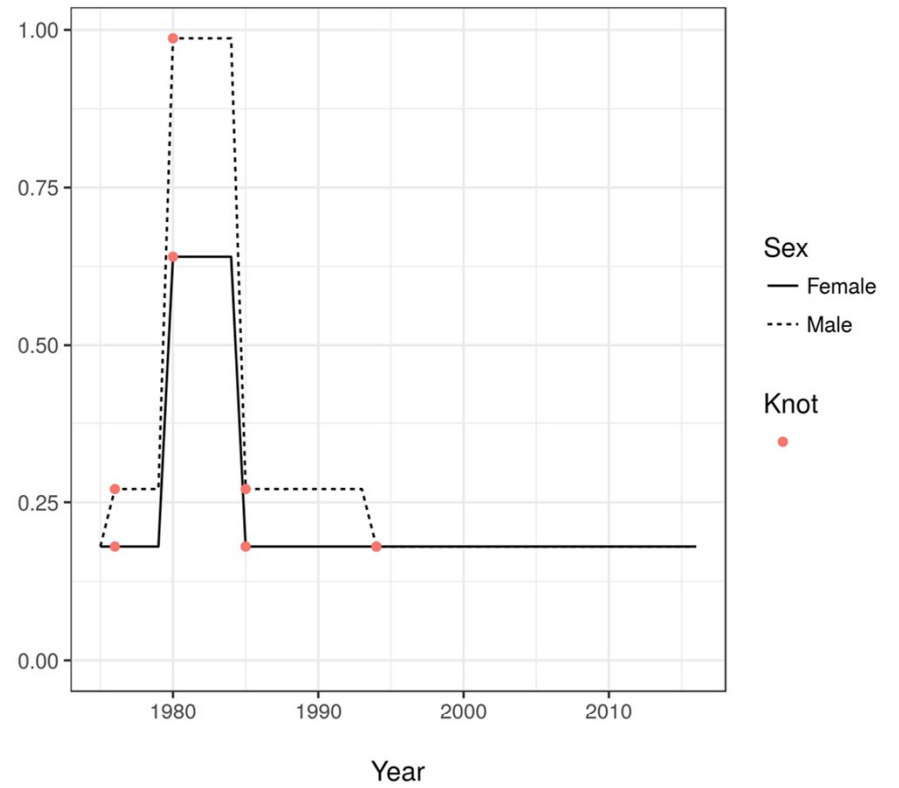
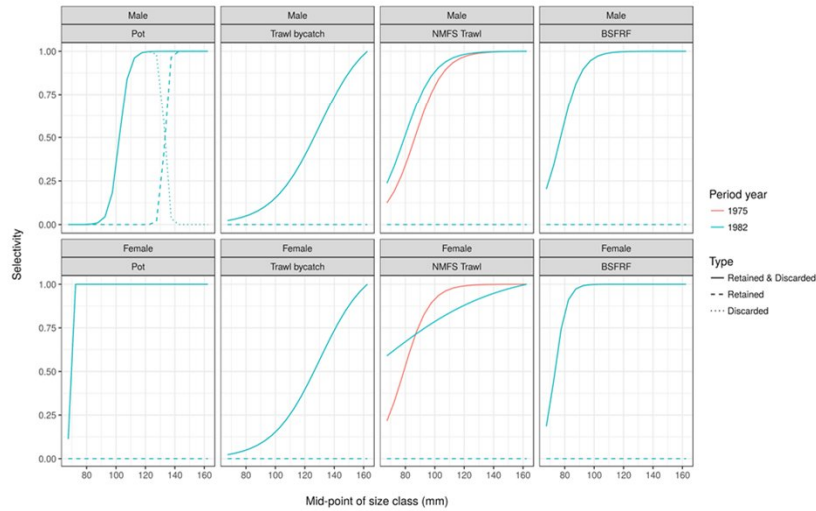
# 3. 2016 Model M



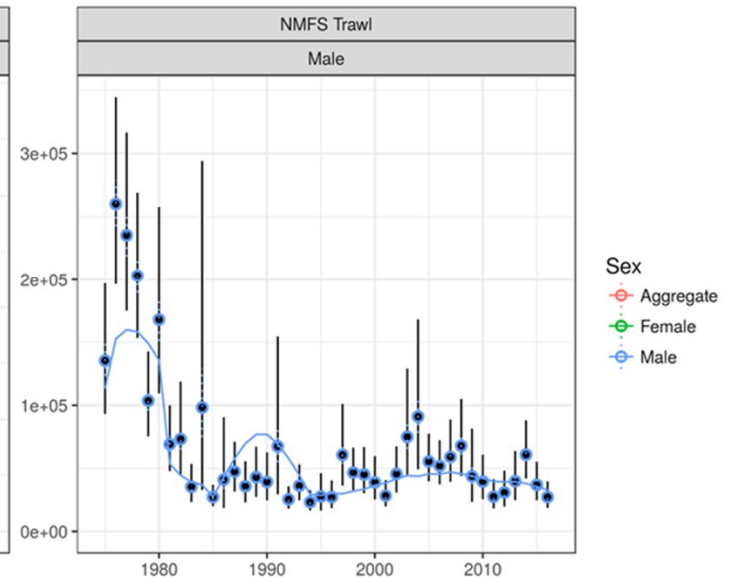
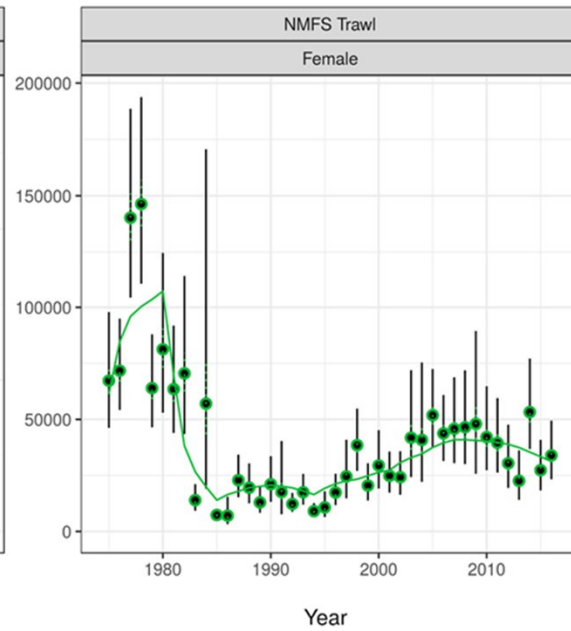
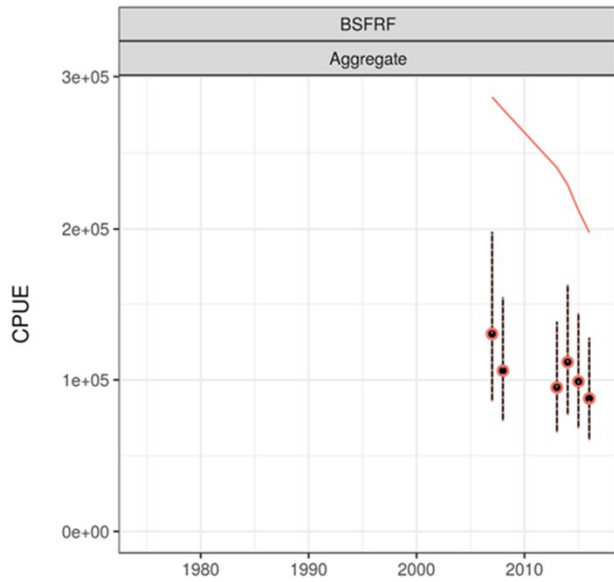
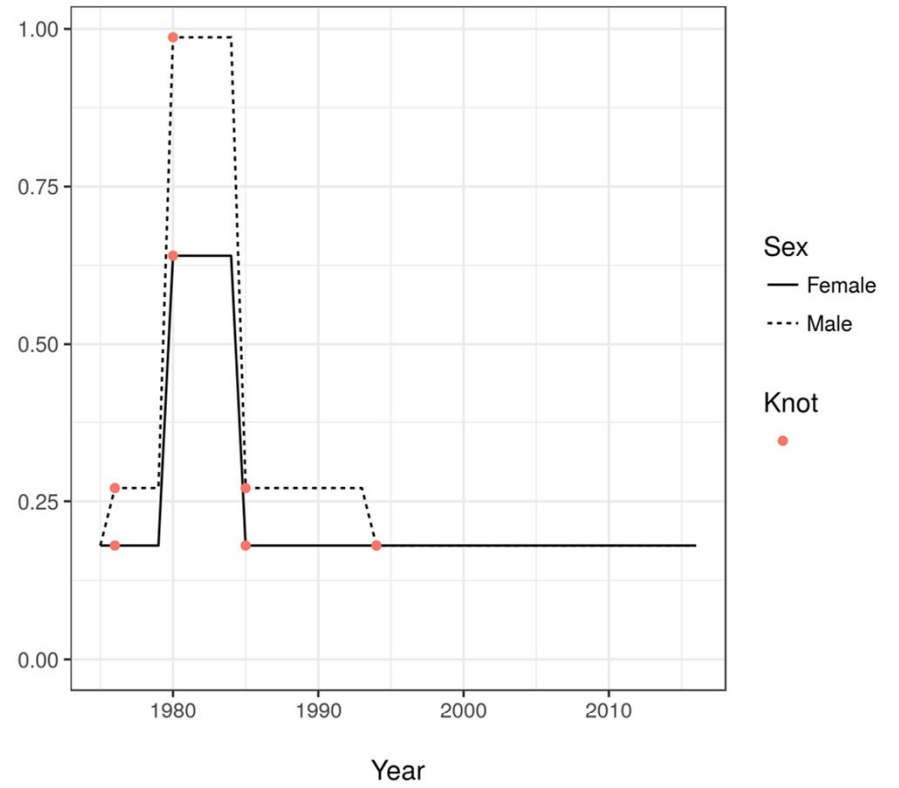
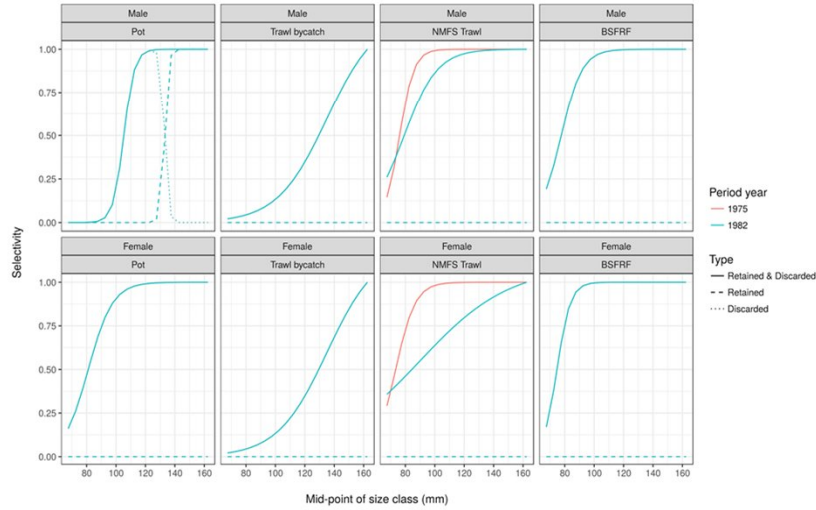
# 3. 2016 Model M



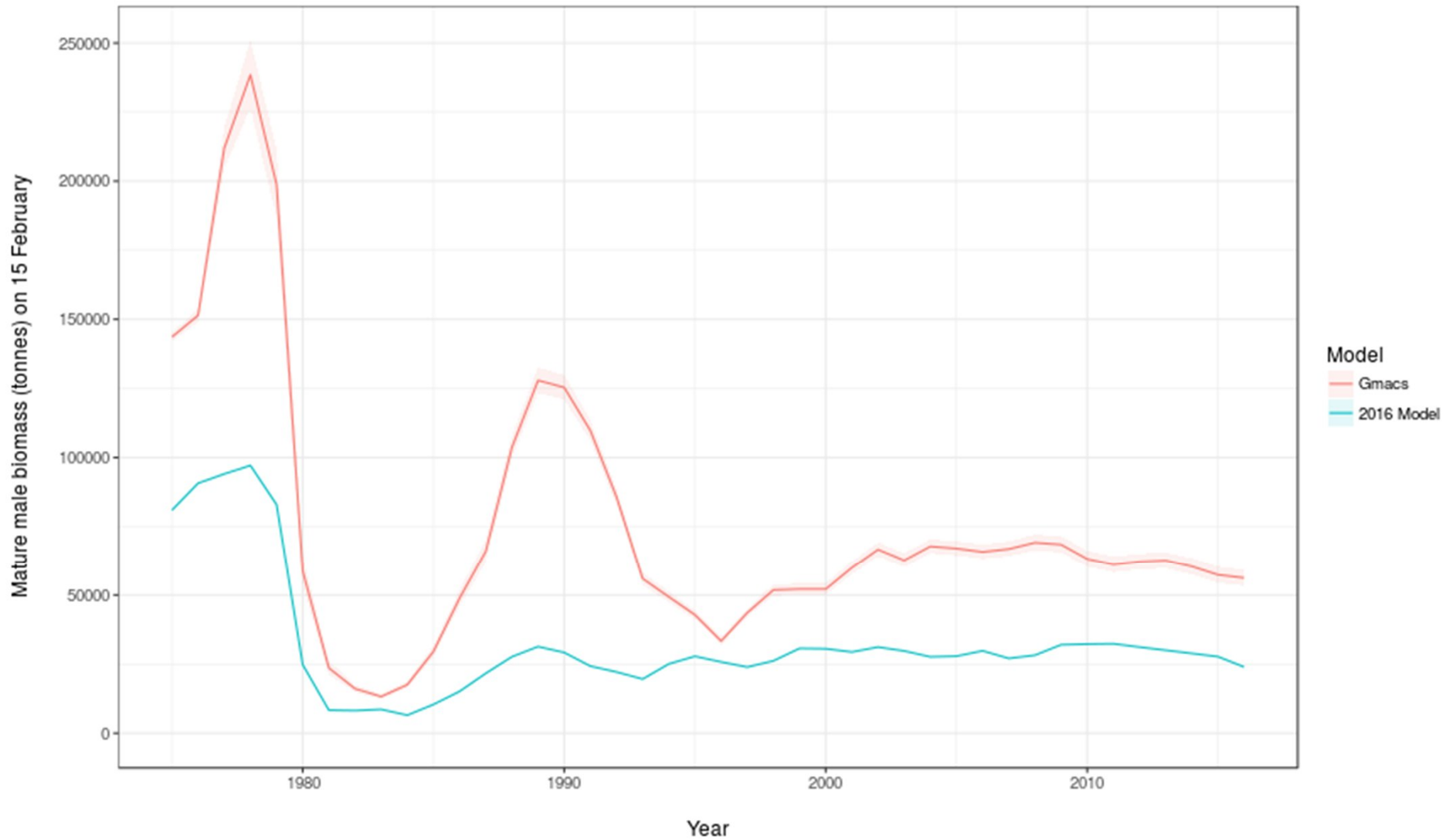
# 4. Estimate BSFRF q



# 5. NMFS lambda = 4

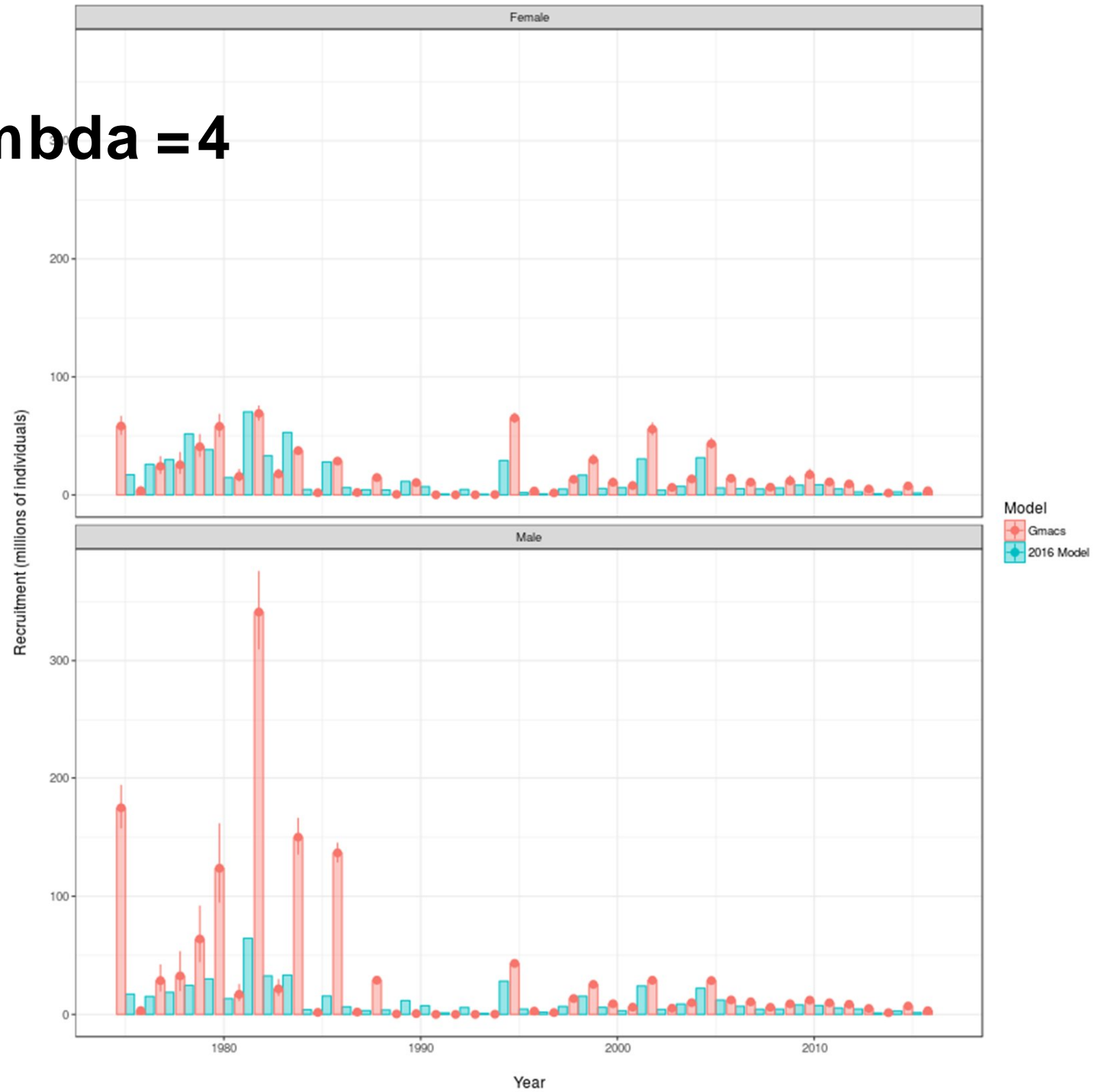


# 5. NMFS $\lambda = 4$

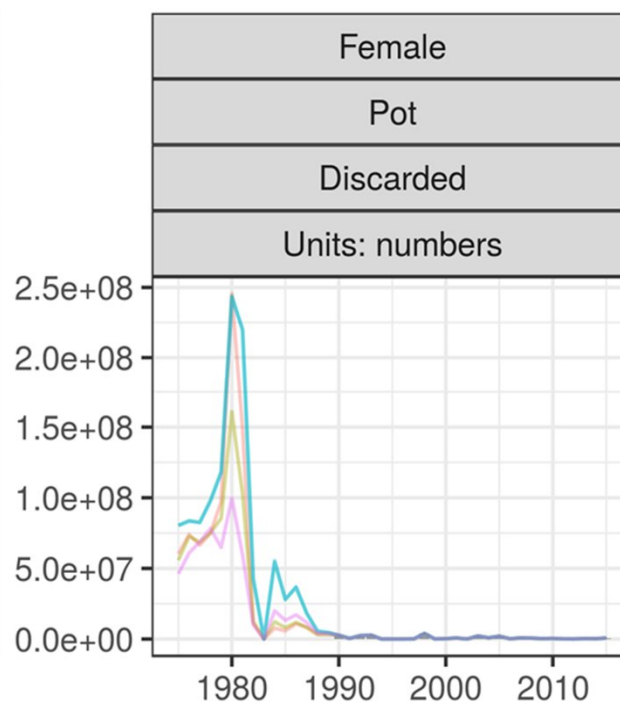
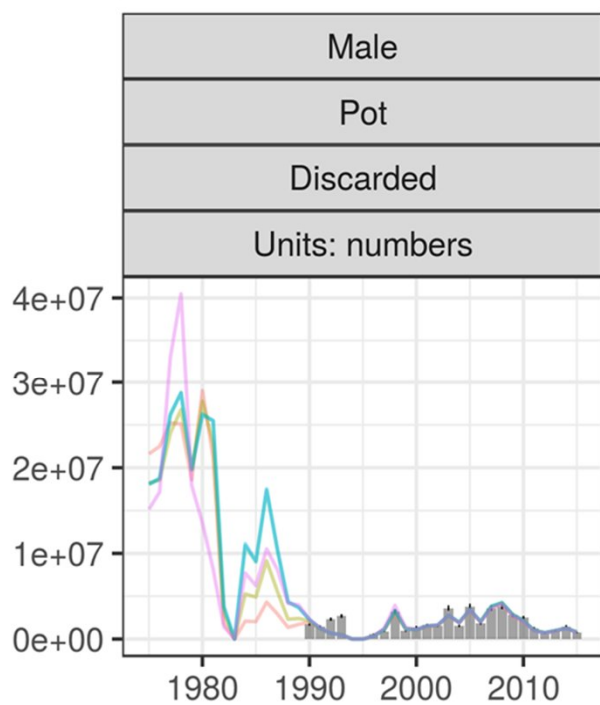
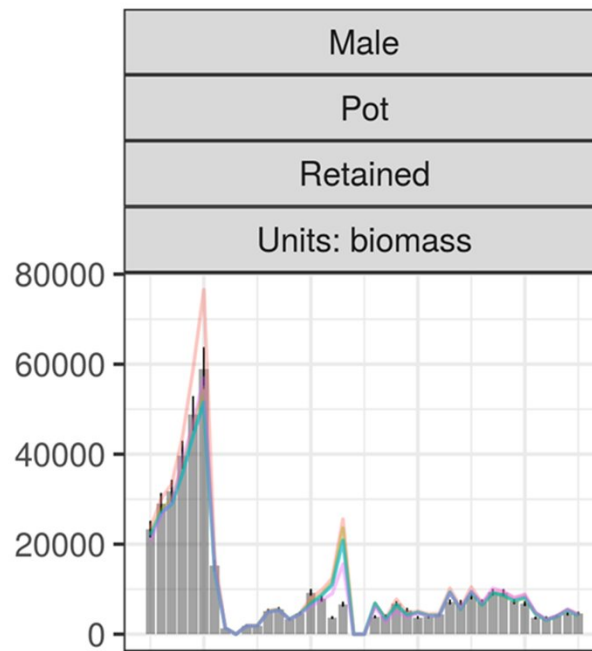
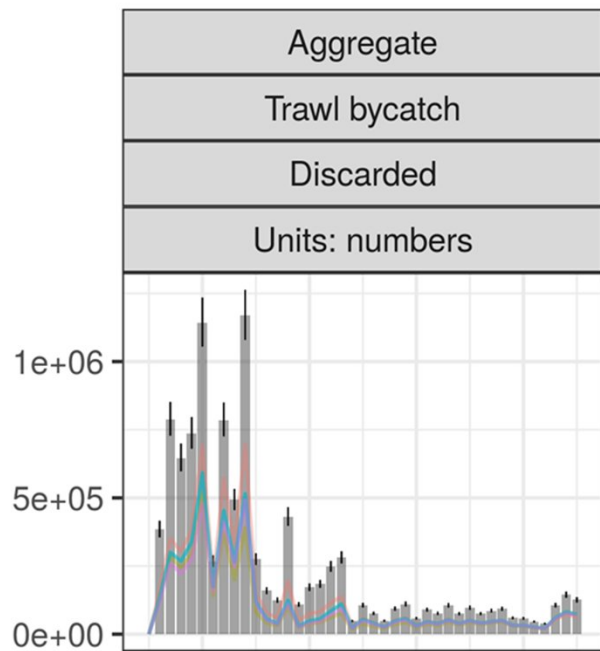




# 5. NMFS lambda = 4



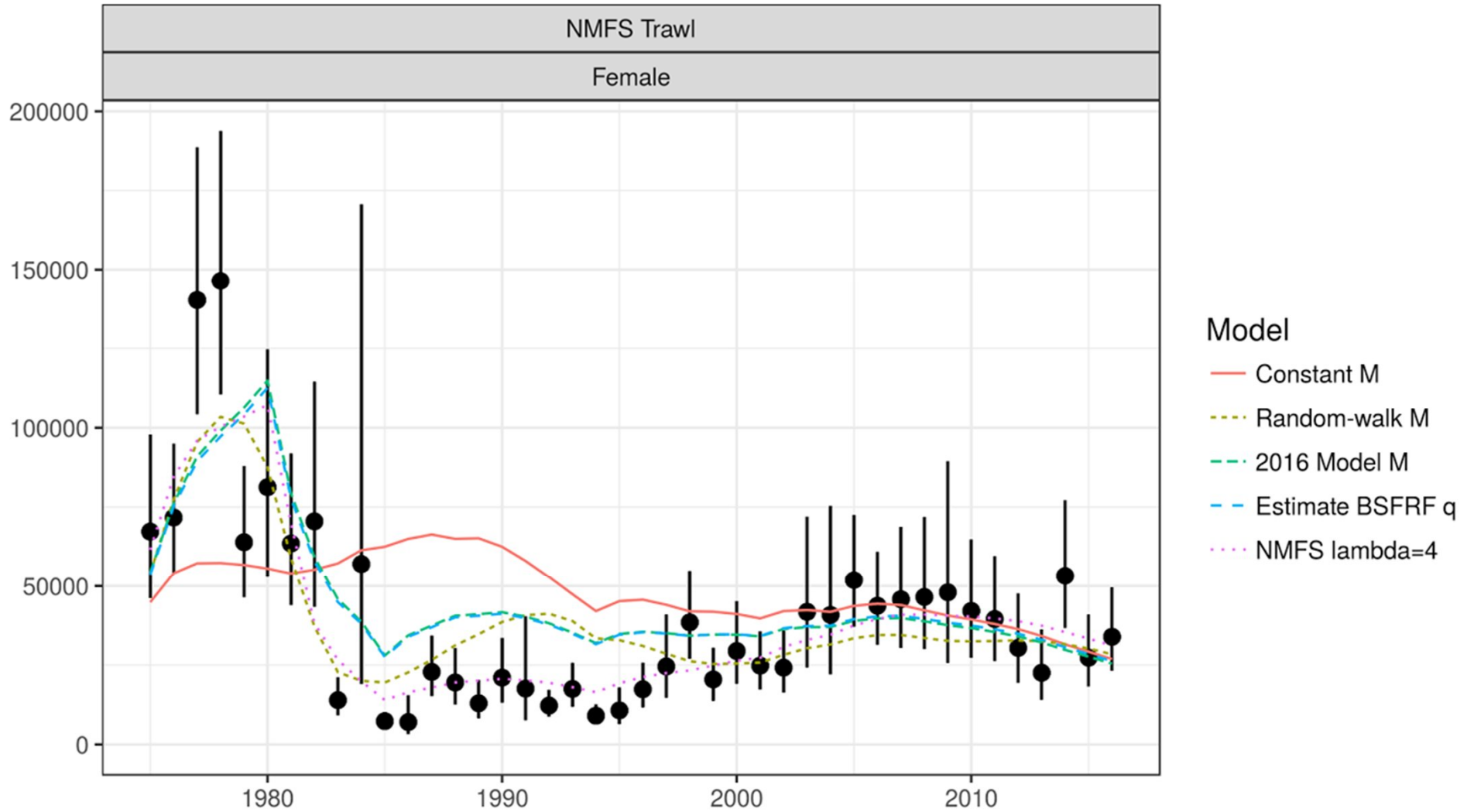
Catch



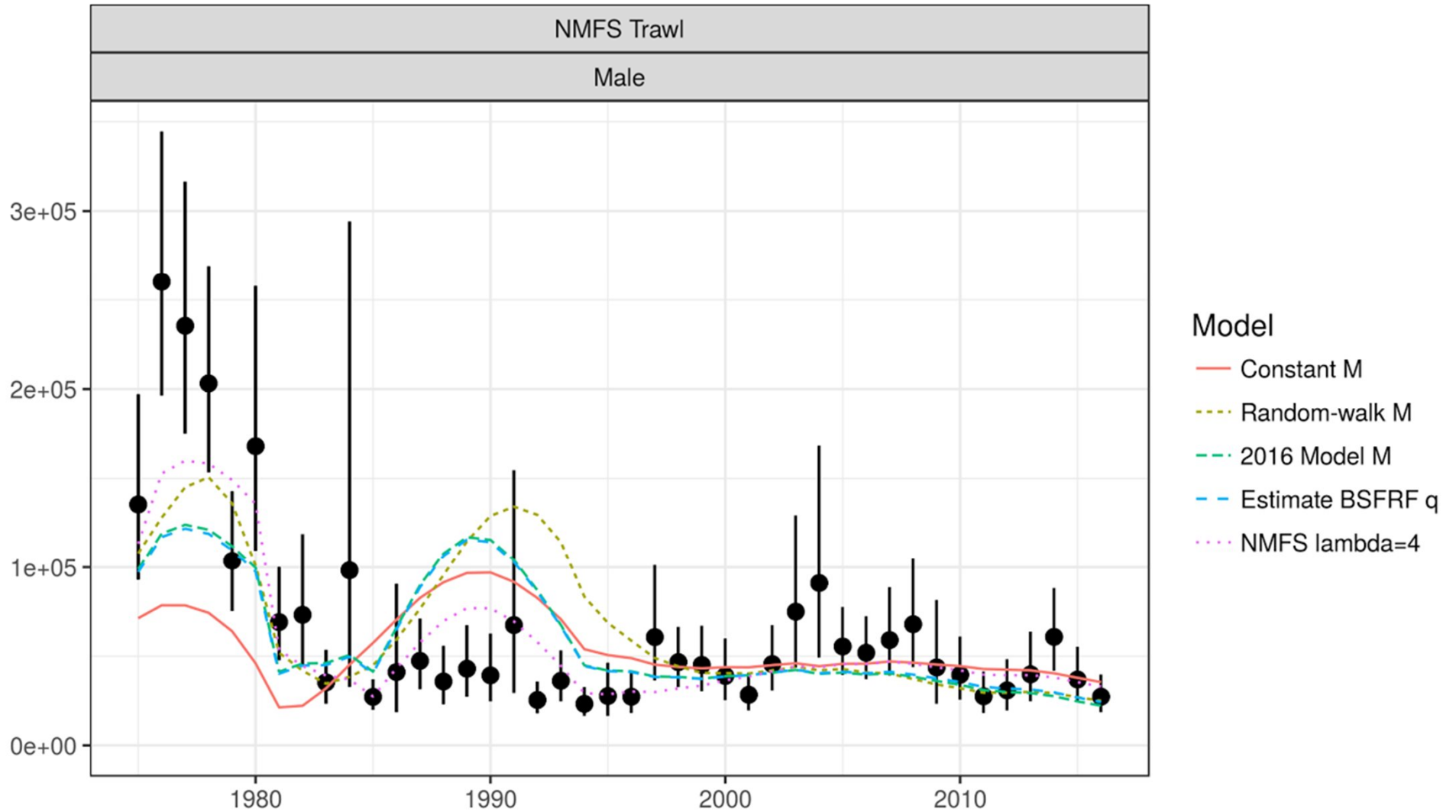
Model

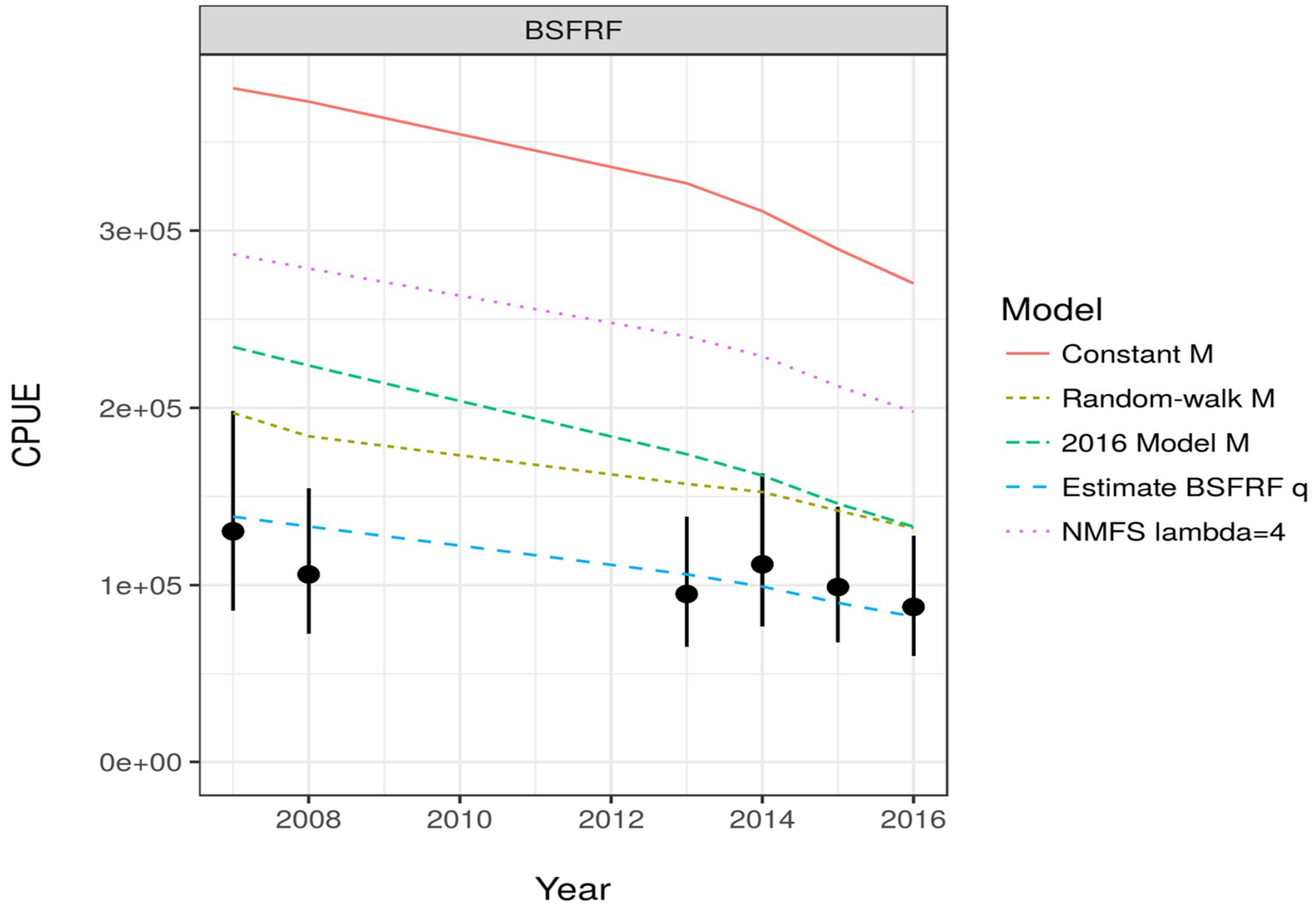
- Constant M
- Random-walk M
- 2016 Model M
- Estimate BSFRF  $q$
- NMFS  $\lambda=4$

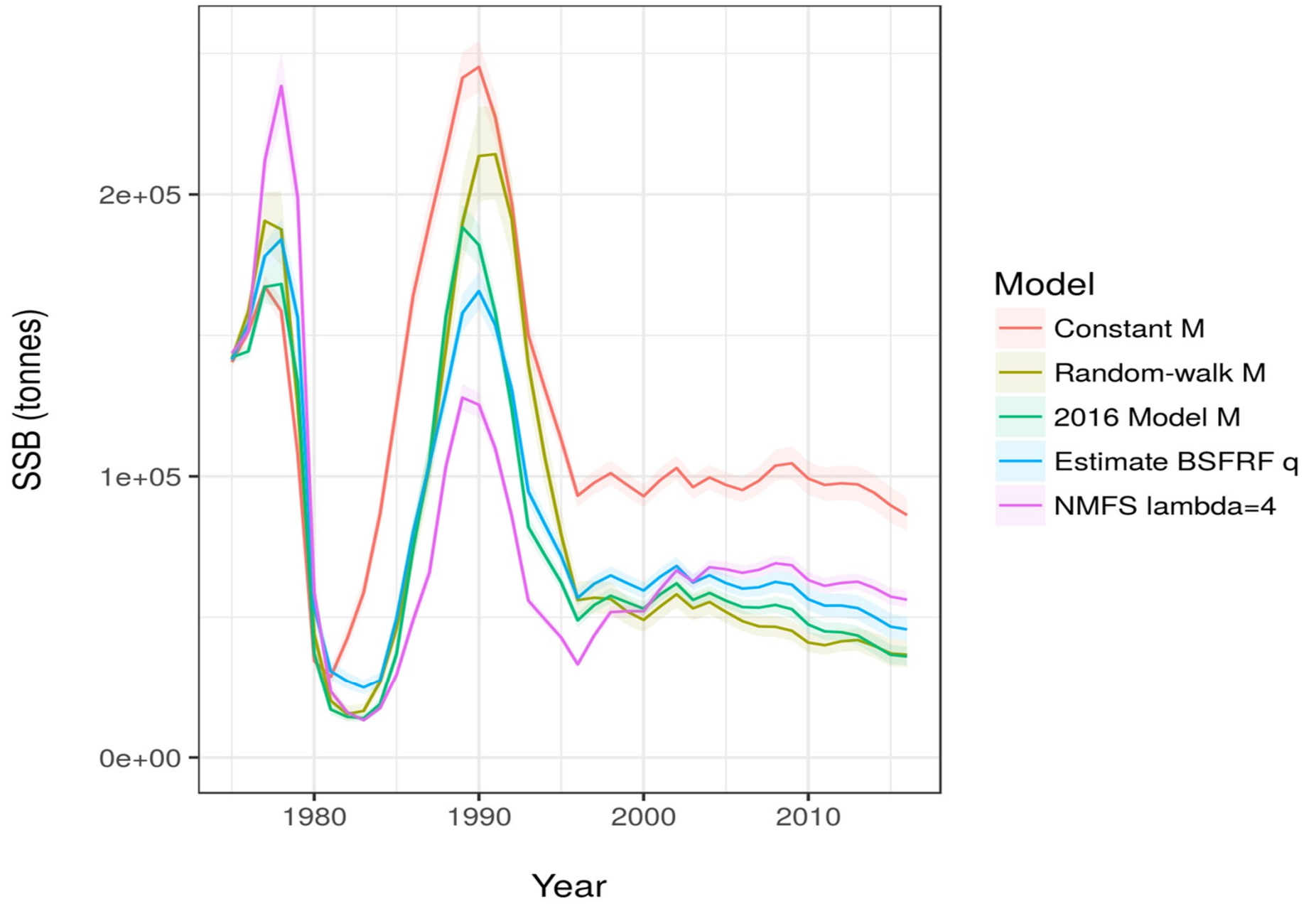
# Model fits (Female)



# Model fits (male)



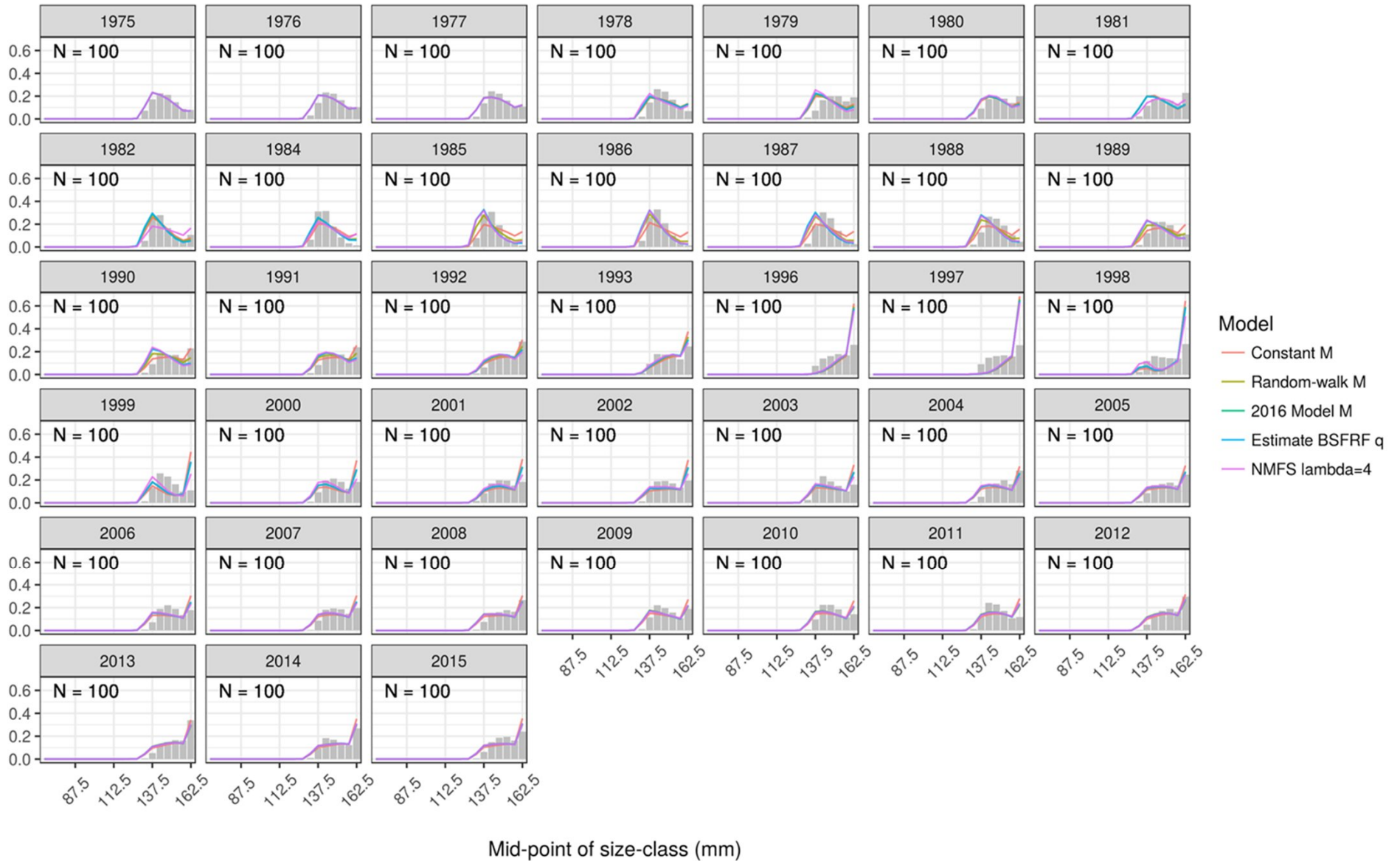






# Retained Males

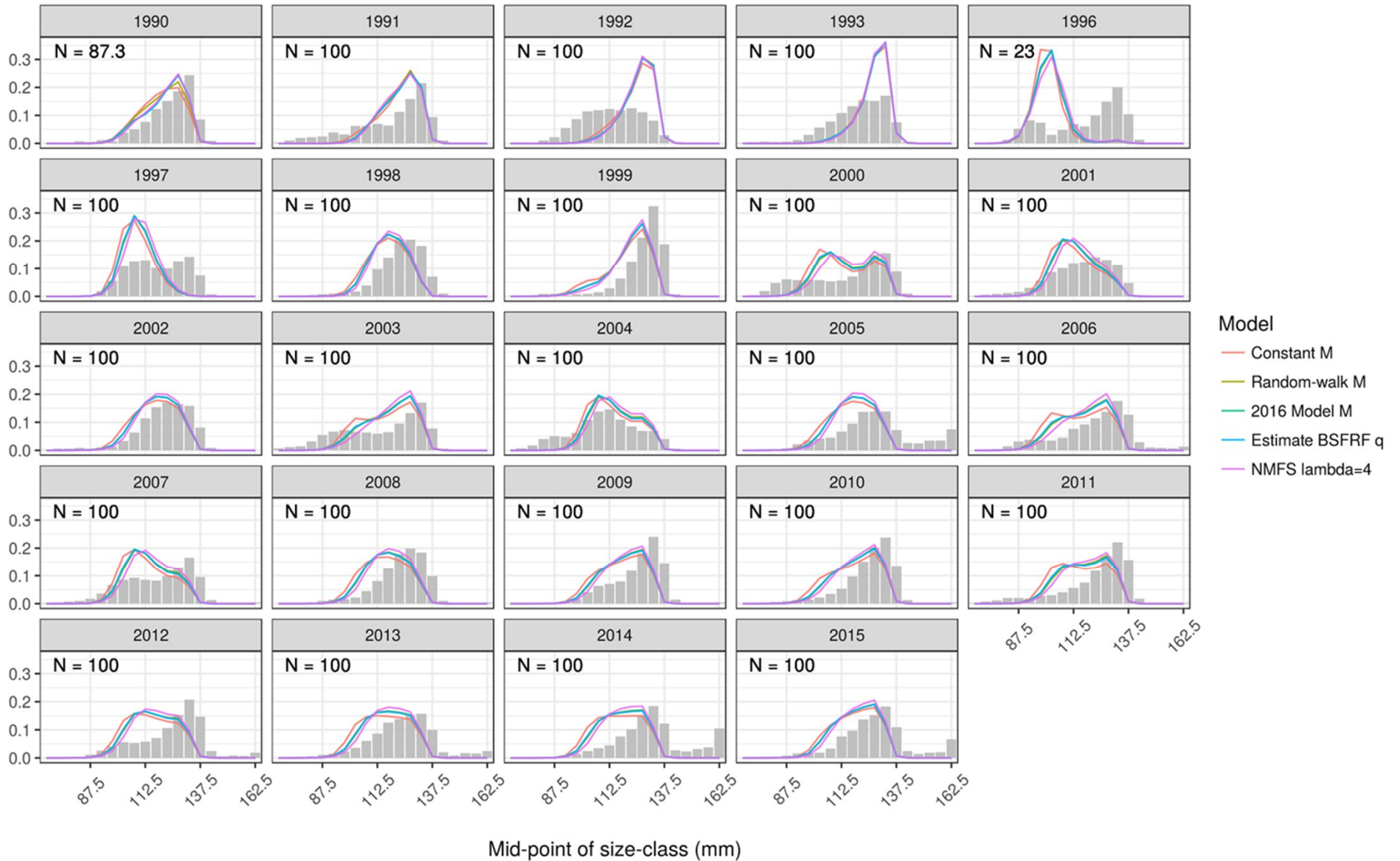
Gear = Pot , Sex = Male , Season = 3





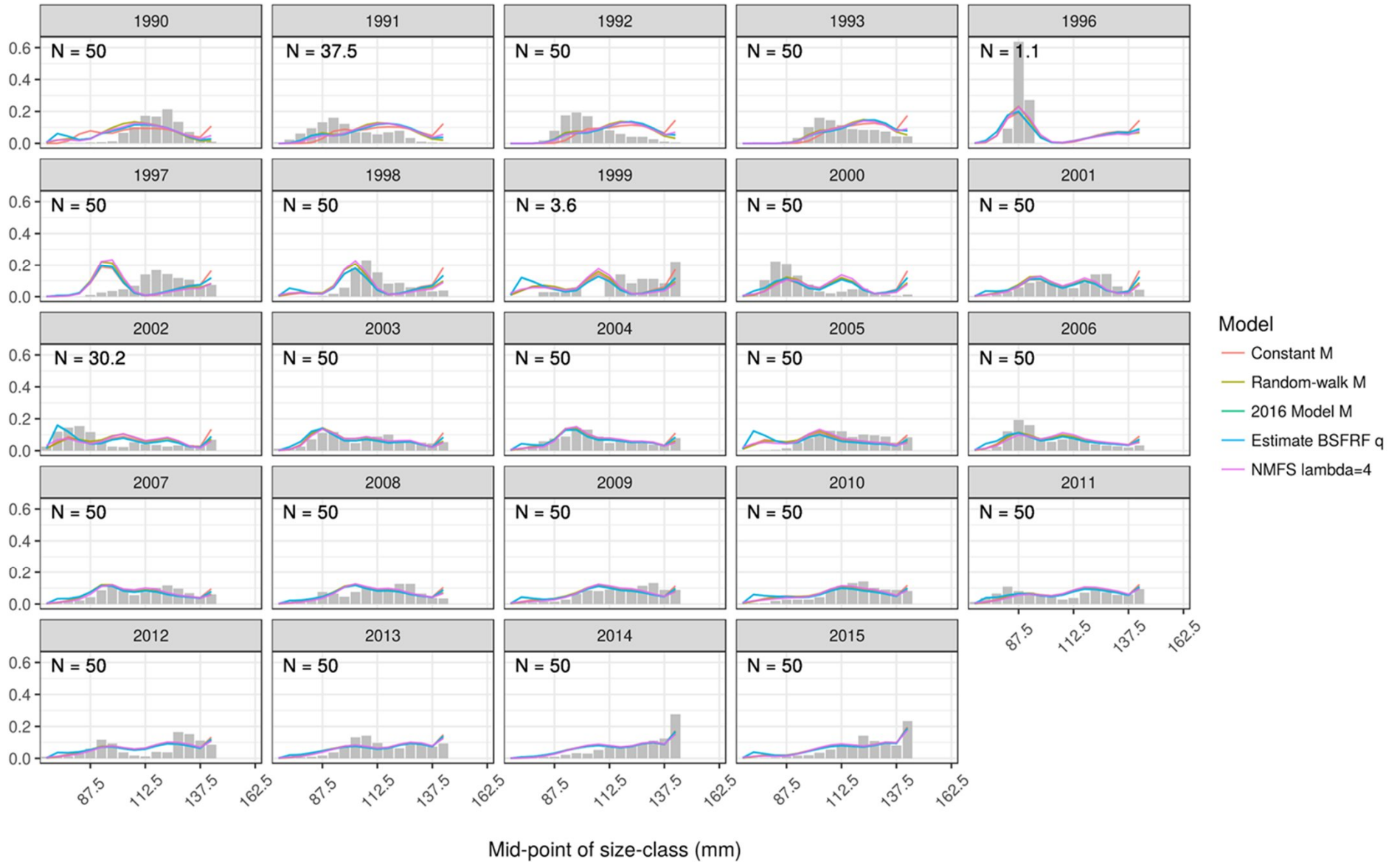
# Discarded Males

Gear = Pot , Sex = Male , Season = 3



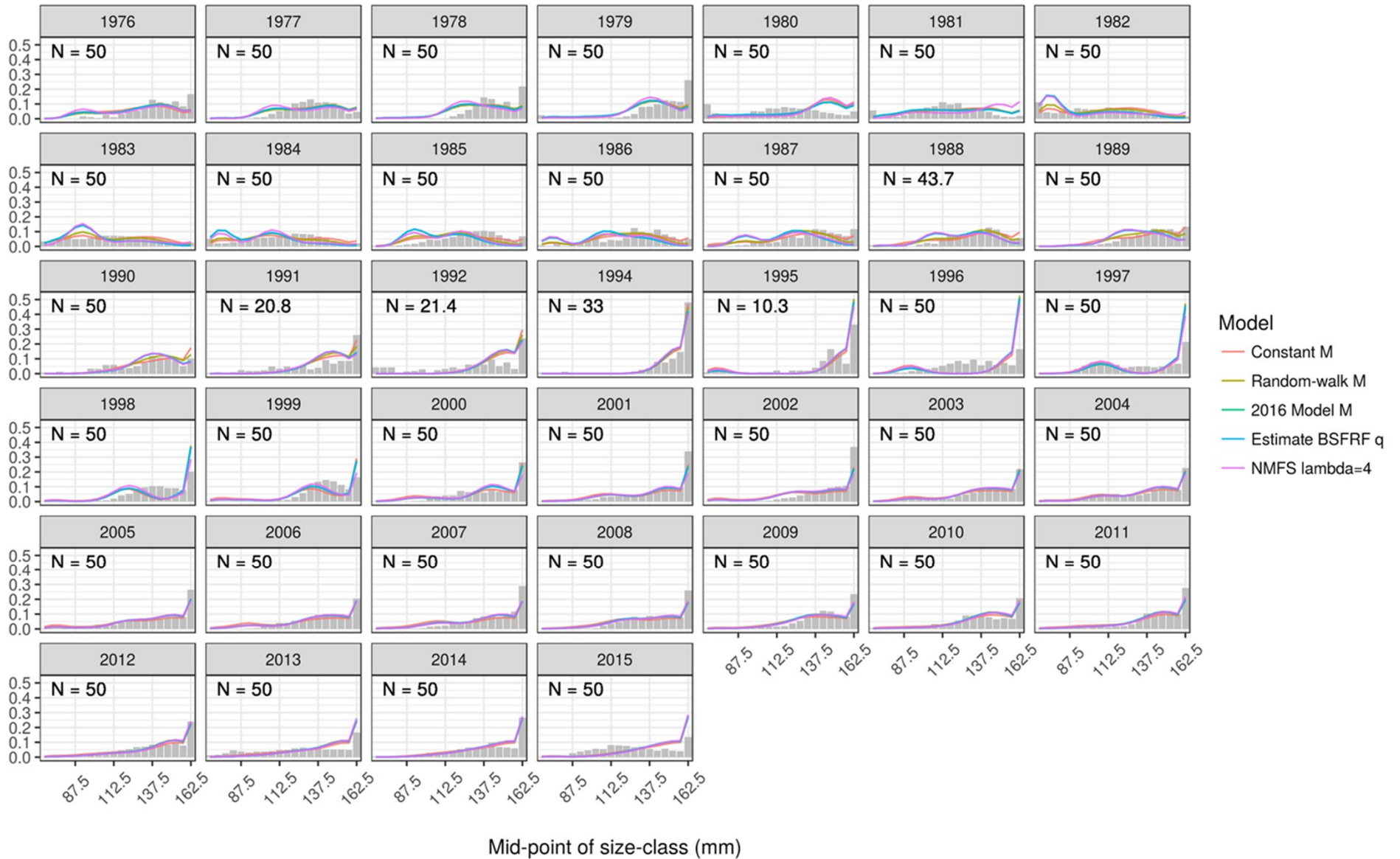
# Discarded Females

Gear = Pot , Sex = Female , Season = 3



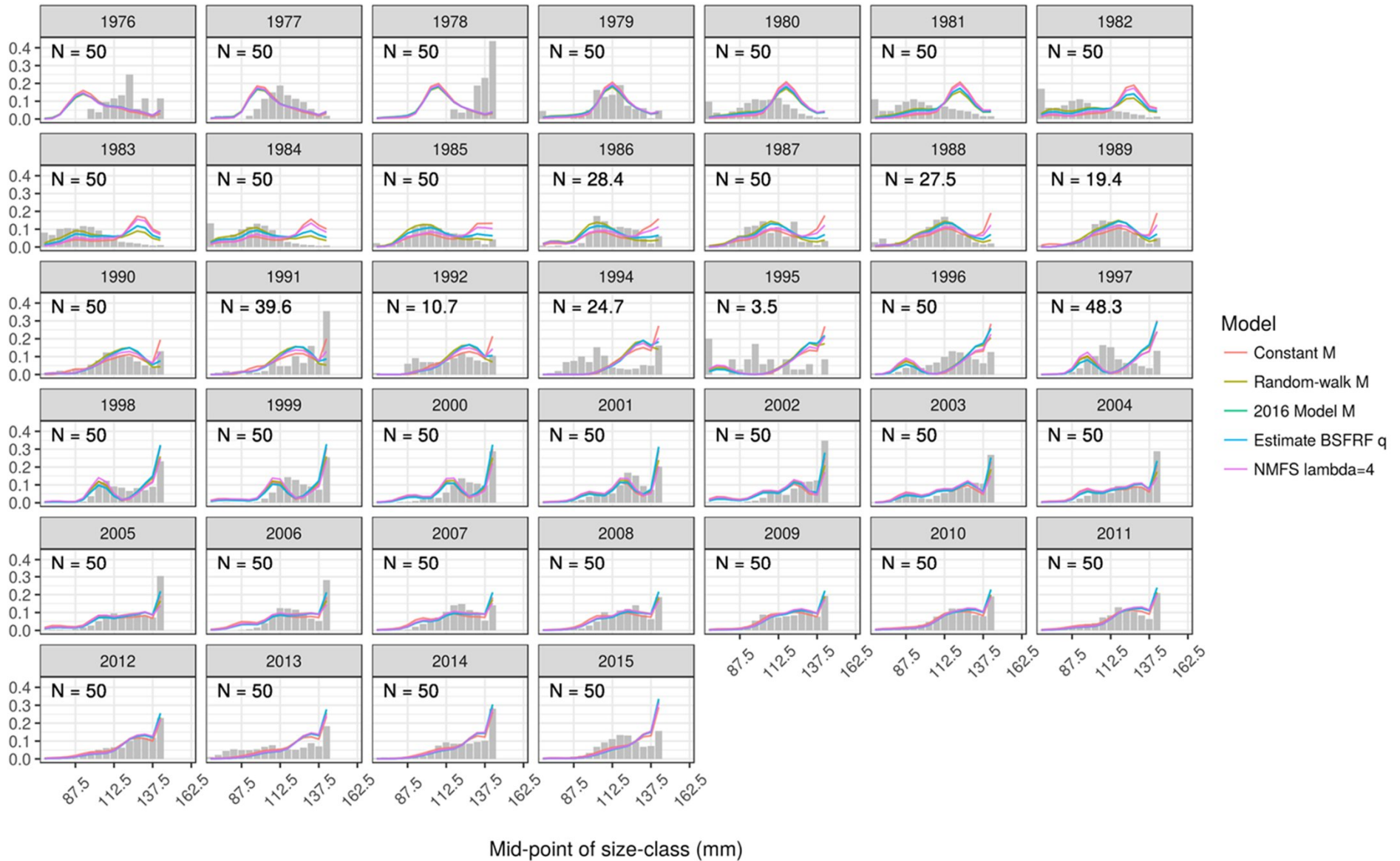
# Trawl Bycatch Males

Gear = Trawl bycatch , Sex = Male , Season = 2



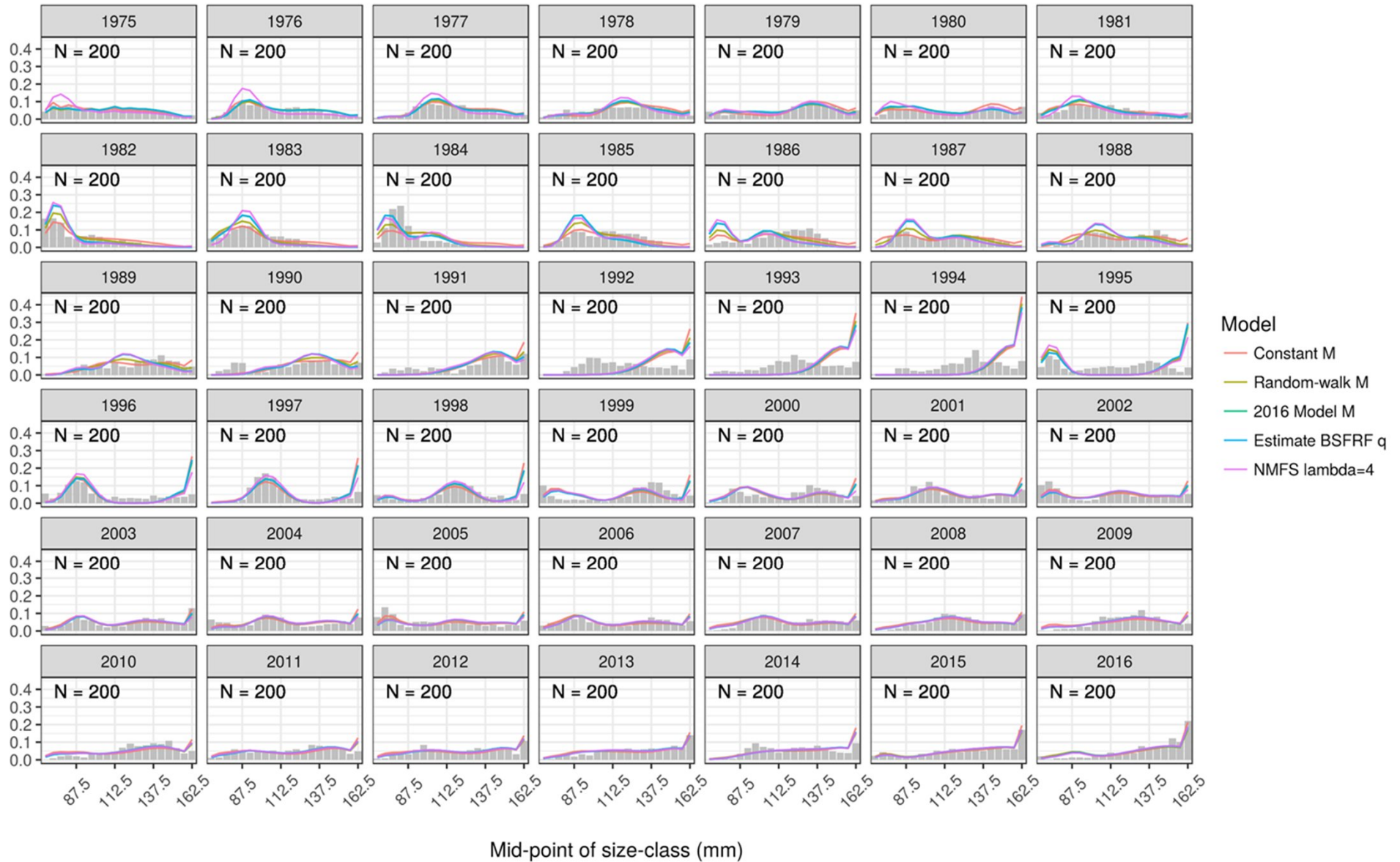
# Trawl Bycatch Females

Gear = Trawl bycatch , Sex = Female , Season = 2



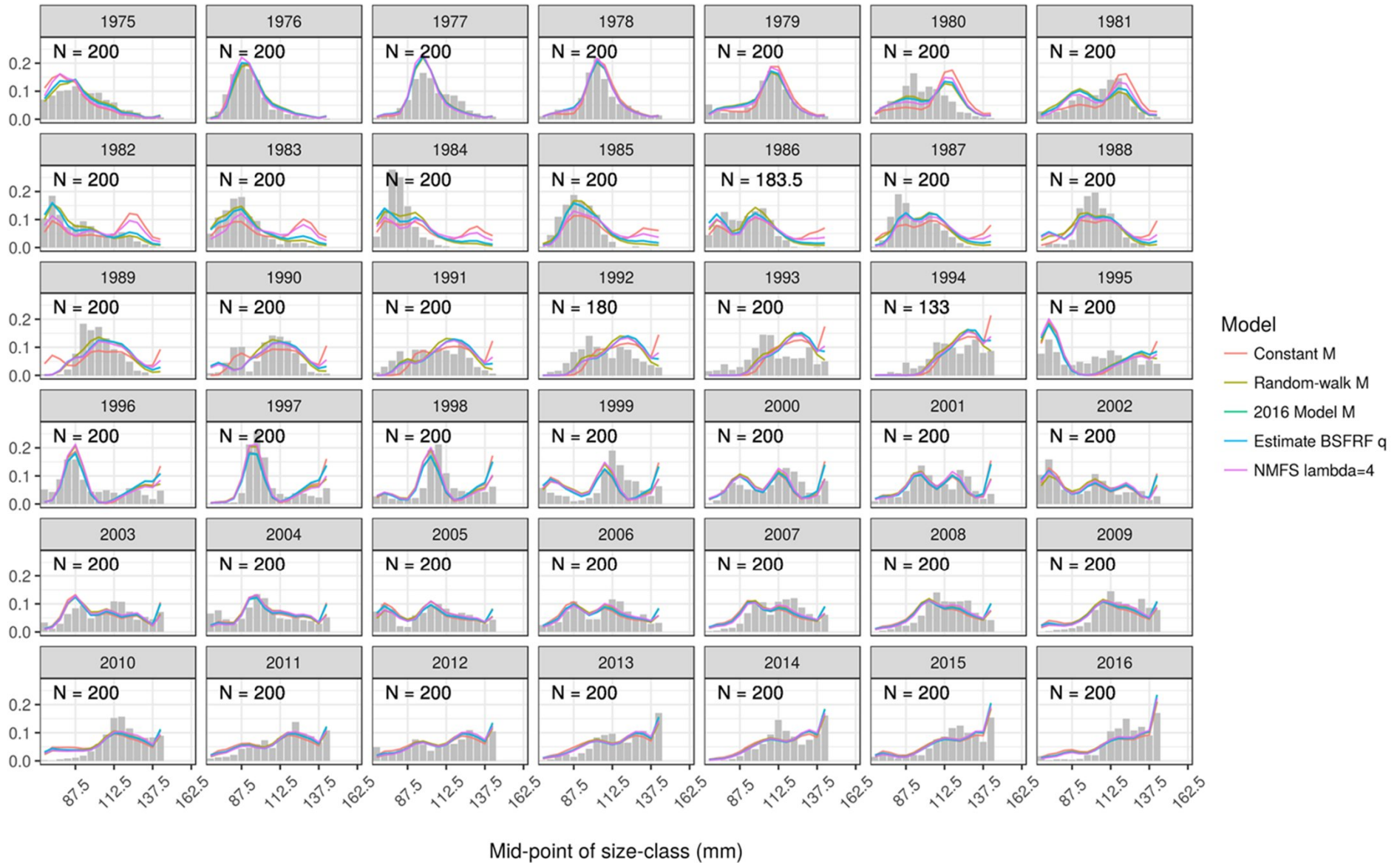
# NMFS Trawl Males

Gear = NMFS Trawl , Sex = Male , Season = 3



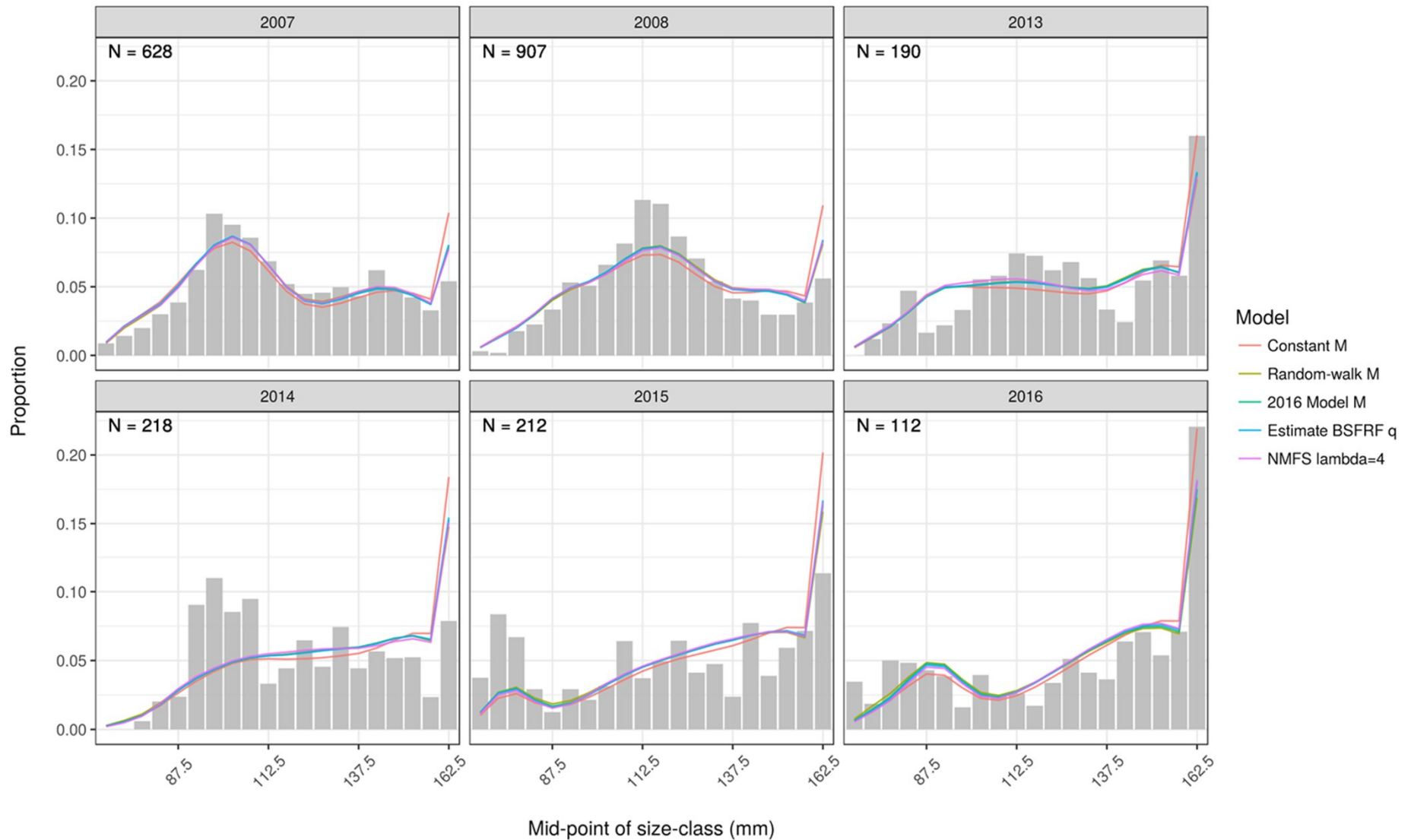
# NMFS Trawl Females

Gear = NMFS Trawl , Sex = Female , Season = 3



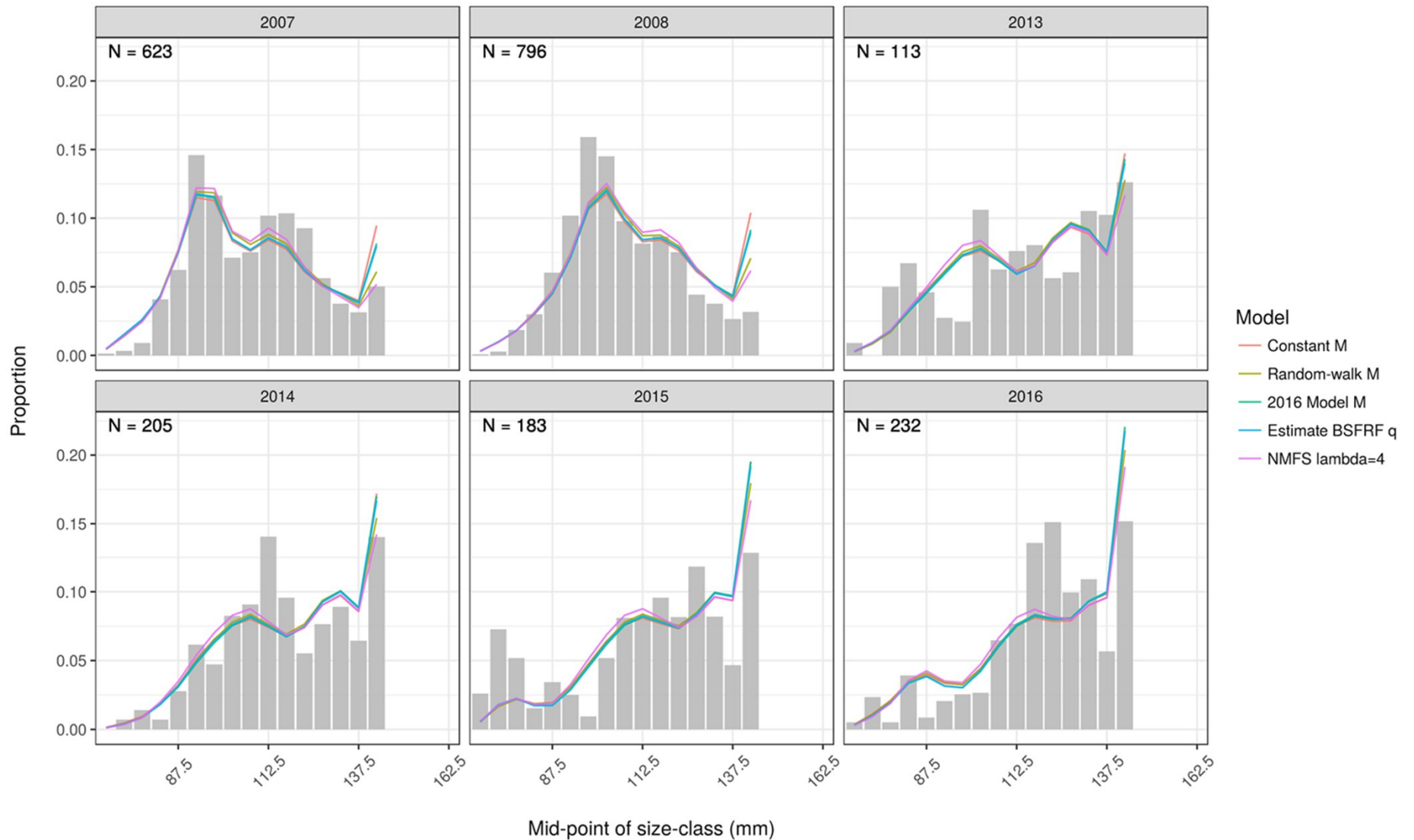
# BSFRF Males

Gear = BSFRF , Sex = Male , Season = 3



# BSFRF Females

Gear = BSFRF , Sex = Female , Season = 3





# To do

1. Need to figure out what is going on with BBRKC initialisation - all of these issues likely stem from this
2. Same goes for growth matrix

This will hopefully result in better survey fits, then:

1. Write-up document

Bristol Bay Red King Crab Stock Assessment 2017

D'Arcy Webber<sup>1</sup>, Jie Zheng<sup>2</sup>, and James Ianelli<sup>3</sup>

<sup>1</sup>Quantifish, darcy@quantifish.co.nz

<sup>2</sup>Alaska Department of Fish and Game, jie.zheng@alaska.gov

<sup>3</sup>NOAA, jim.ianelli@noaa.gov

January 2017

## Executive Summary

1. **Stock:** Red king crab (RKC), *Paralithodes camtschaticus*, in Bristol Bay, Alaska.
2. **Catches:** Peak historical harvest was 4288 tonnes (9.454 million pounds) in 1983/84<sup>1</sup>. The fishery was closed for 10 years after the stock was declared overfished in 1999. Fishing resumed in 2009/10 with a fishery-reported retained catch of 209 tonnes (0.461 million pounds), less than half the 529.3 tonne (1.167 million pound) TAC. Following three more years of modest harvests supported by a fishery catch per unit effort (CPUE) of around 10 crab per pot lift, the fishery was again closed in 2013/14