

Today's Specials

Pollock

Cod

Flounder



# Estimating Predation Mortality with a Three-Species Model in the Gulf of Alaska

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# General Approach

**Construct a set of simplified single-species models and connect them through predation.**

## ■ **SSM: Single-species models**

- Designed to match stock assessment output
- $Z = F + M$

## ■ **MSM: Multispecies model**

- Calculates predation from gut data assumed to be measured without error
- $Z = F + M + P$

## ■ **MSASA: Multispecies Age-Structured Assessment**

- Estimates predation from a series of model parameters
- $Z = F + M + P$

# Three species with close predator-prey links

Pacific Cod (*Gadus macrocephalus*)

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Arrowtooth Flounder  
(*Atheresthes stomias*)



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## Work forward from Age 1 cohort

$$N_{i,a+1,t+1} = N_{i,a,t} e^{(-Z)}$$

- No recruitment function: Age 1 is a model parameter.

- **Deconstruct  $Z$  into:**
  - **Fishing mortality  $F$**
  - **Predation mortality  $P$**
  - **Residual natural mortality  $M$**

$$N_{i,a+1,t+1} = N_{i,a,t} e^{(-M - F - P_1 - P_2 \dots P_n)}$$

**The Multispecies Model (MSM) and the Multispecies Age-Structured Assessment (MSASA) are simply extensions of the SSM in which  $Z = F + M + P$**



# Natural Mortality

- **SSM**: Set to values used in the stock assessments
  - Ages 1 and 2 for prey species set higher to mimic  $P$

	SSM
ATF 1	0.45
ATF 2	0.4
ATF 3+	0.24
COD	0.37
PLK 1	0.9
PLK 2	0.8
PLK 3+	0.3

# Natural Mortality

- **MSM / MSASA**: Set all ages to stock assessment values.

	SSM
ATF 1	0.45
ATF 2	0.4
<b>ATF 3+</b>	<b>0.24</b>
<b>COD</b>	<b>0.37</b>
PLK 1	0.9
PLK 2	0.8
<b>PLK 3+</b>	<b>0.3</b>

# Modeling predation

$$P_{i,a,t} = \frac{1}{N_{i,a,t} W_{i,a}} \sum_j \sum_b I_{j,b} N_{j,b,t} \frac{\phi_{i,a,j,b,t}}{\phi_{j,b,t}}$$

$i$  = prey species  
 $j$  = predator species  
 $a$  = prey age  
 $b$  = predator age

Annual Ingestion  $\times$  Predator abundance

Total ingestion by predator  $j$  \* Proportion of the ingested food that is prey  $i$ , age  $a$

$\Sigma$  Total amount of prey  $i$  consumed by predator  $j$

$$P_{i,a,t} = \frac{\text{Total amount of prey } i \text{ consumed by all predators}}{\text{Biomass of prey } i}$$

# Predation for the MSM

$$P_{i,a,t} = \frac{1}{N_{i,a,t} W_{i,a}} \sum_j \sum_b I_{j,b} N_{j,b,t} \frac{\phi_{i,a,j,b,t}}{\phi_{j,b,t}}$$


**Proportion of ingested food for predator  $j,b$  composed of prey  $i,a$  taken directly from gut studies.**

# Predation for the MSASA

## Estimating stomach contents

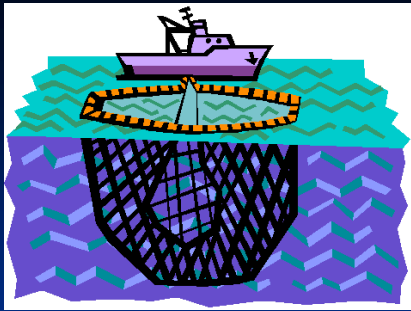
**SIZE-PREFERENCE:** Defined by the gamma function

**SUITABILITY:** size-preference \* prey vulnerability

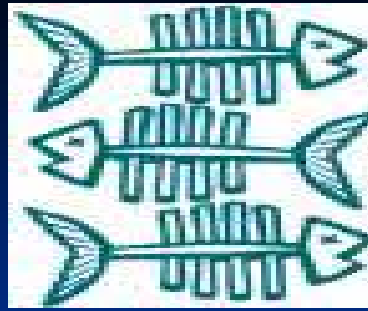
• Scaled to 1 to create a distribution of gut contents for each predator

**AVAILABILITY:** suitability \* prey biomass

$$\text{Stomach} = \frac{\text{Biomass of prey } i,a \text{ available to predator } j,b}{\text{Total food available to predator } j,b}$$



**Age-Specific  
Fishing Mortality**



**Residual Natural  
Mortality**



**Age-Specific  
Predation Mortality**



**YEAR ONE**



**Age 1 Abundance**



**YEAR TWO**



# Likelihood Components

- Catch-at-age
- Survey-at-age
- Total Annual Catch
- Estimated survey biomass

Additional MSASA component:

- Gut data

# Results

**The SSM is a valid baseline for evaluating predation.**

**Matches stock assessment ranges for:**

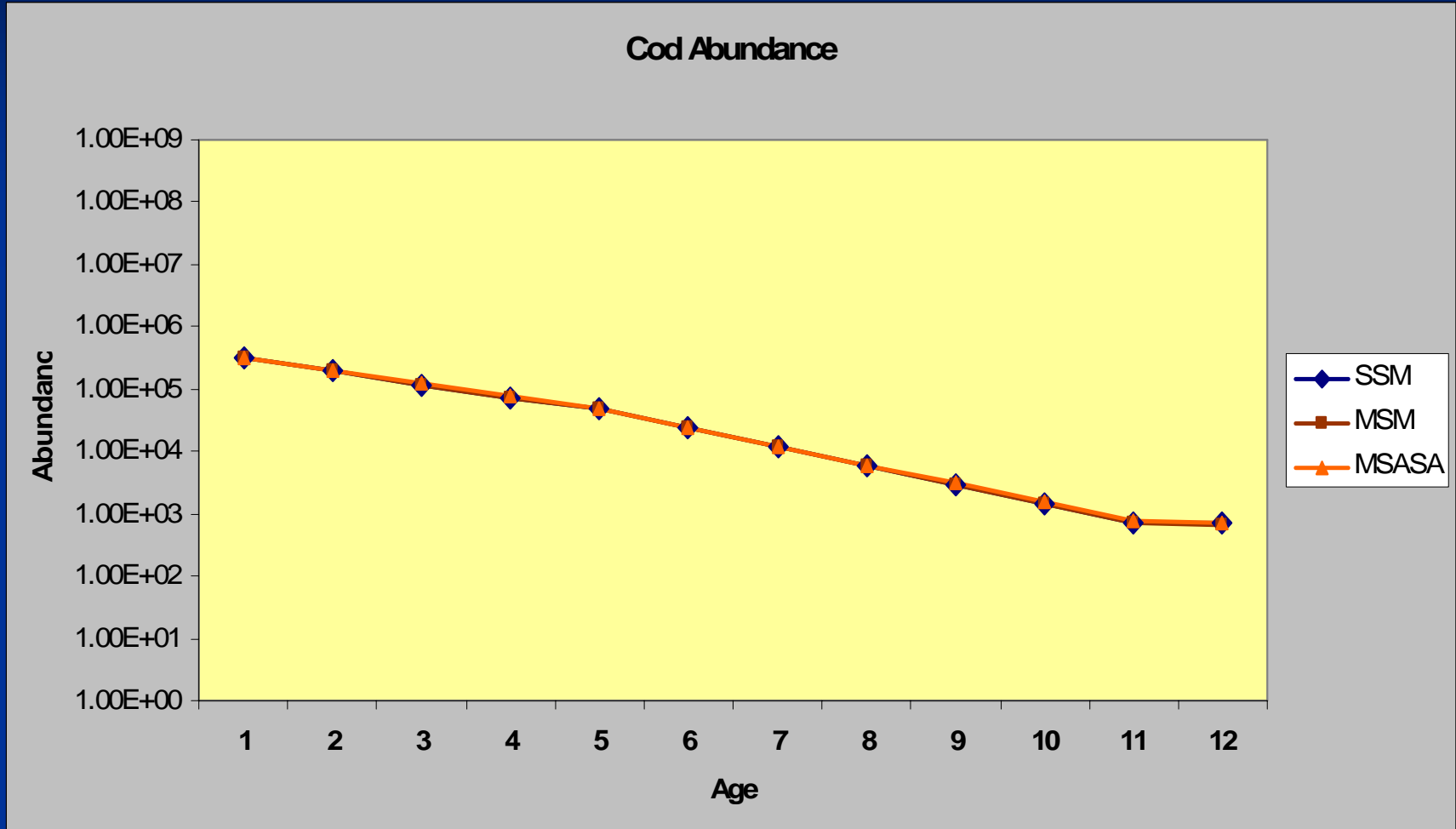
- **Catch and Survey data**
- **Values for  $F$  and  $Q$**
- **Gear selectivity patterns**
- **Estimates of abundance**

**We assume that stock assessment estimates of biomass, abundance,  $Q$ ,  $F$ , and gear selectivities are essentially correct.**



# Results

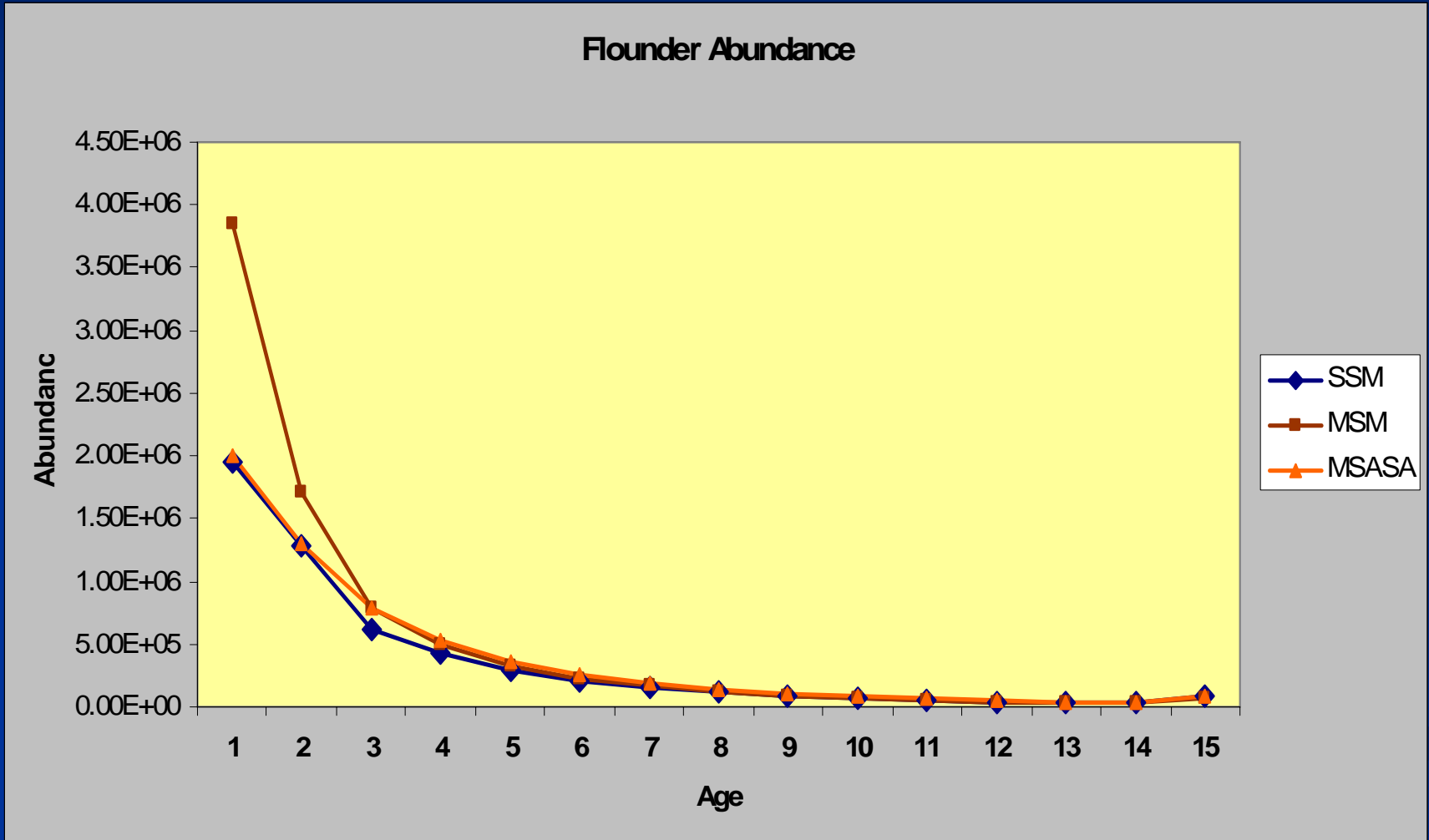
## Abundance



No predation = no change in abundance from SSM

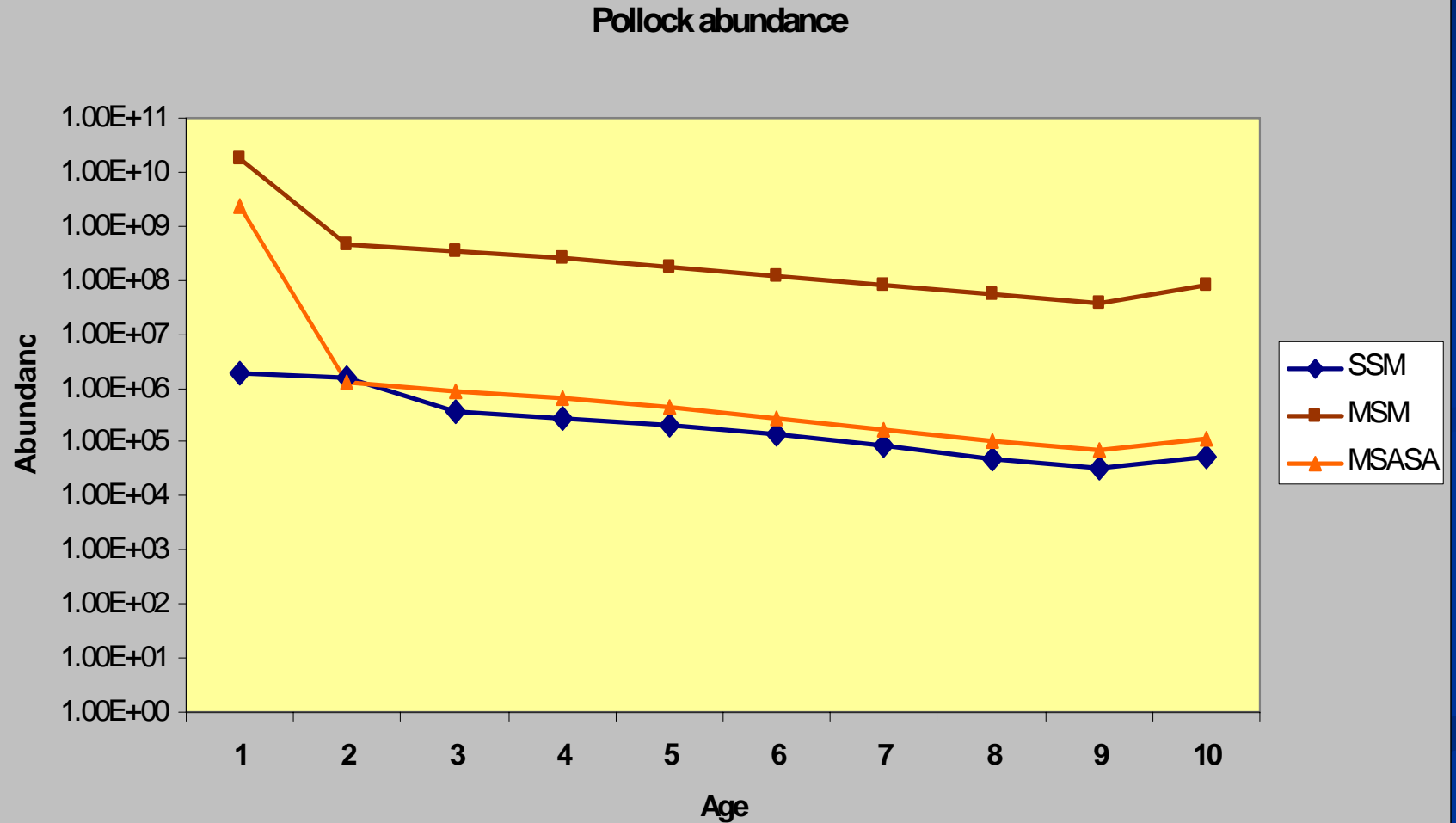
# Results

## Abundance



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## Abundance



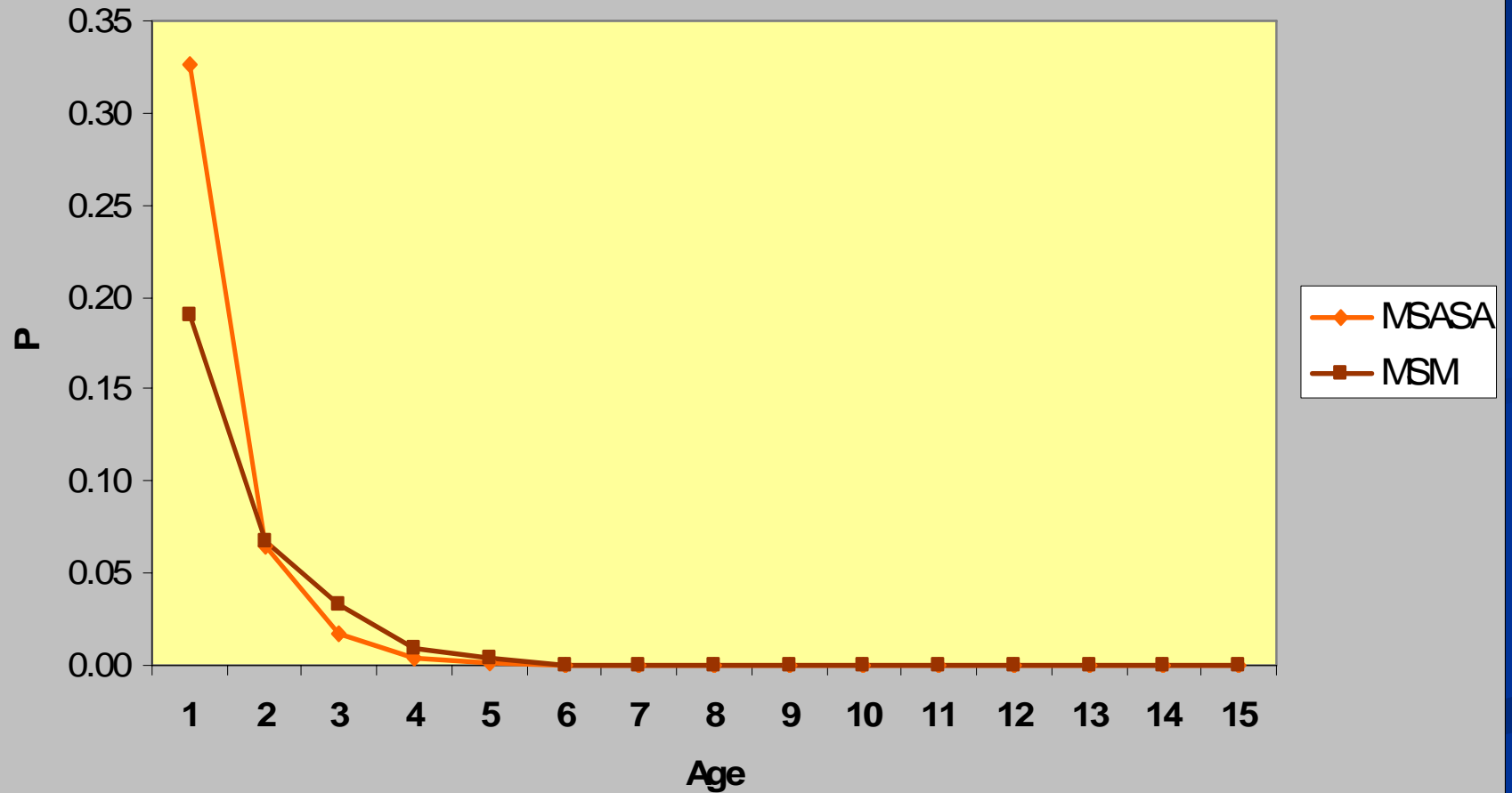
# Predation Mortality

P on Pollock



# Predation Mortality

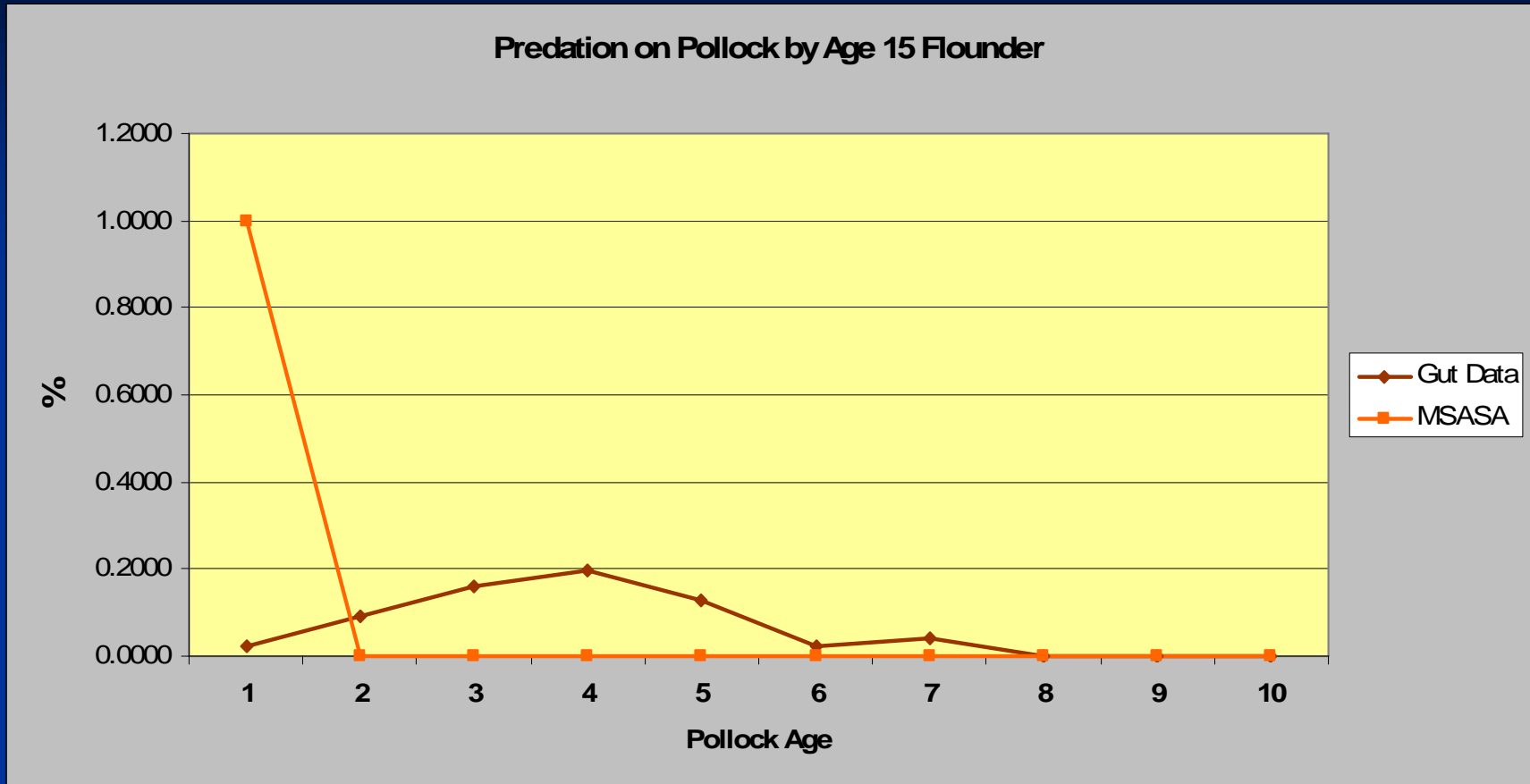
P on Flounder



# Catch and Survey Factors

	<b>SSM</b>	<b>MSM</b>	<b>MSASA</b>
ATF Q	1	1	0.88
PLK Q	0.75	0.0009	0.46
ATF F	N/A	N/A	N/A
PLK F	0.15	0.0002	0.1

# Issues: Gut Content



- 1. Weak forcing from gut data: too many zeroes**
- 2. Large majority of predation on Age 1 Pollock skews model predictions**

# Issues: Natural Mortality

Traditional single-species models have included predation under the umbrella of  $M$ .

By separating  $P$  from  $M$ , values for  $M$  should decrease.

When set as model parameter,  $M$  increases with the addition of predation. Why?



# In Sum

## OBSERVED

- Predation mortality is separable from natural mortality
- Predation mortality is estimable
- The MSASA is both computationally workable and biologically sound

## REQUIRED

- Improved assessment of gut data and its integration into model function
- Improved understanding of relationship between  $M$ ,  $P$ , and cohort dynamics

**What's going on down there??**

