Can we estimate the abundance of yellowfin tuna?

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How we estimate the abundance of yellowfin tuna in the EPO?

Current method:

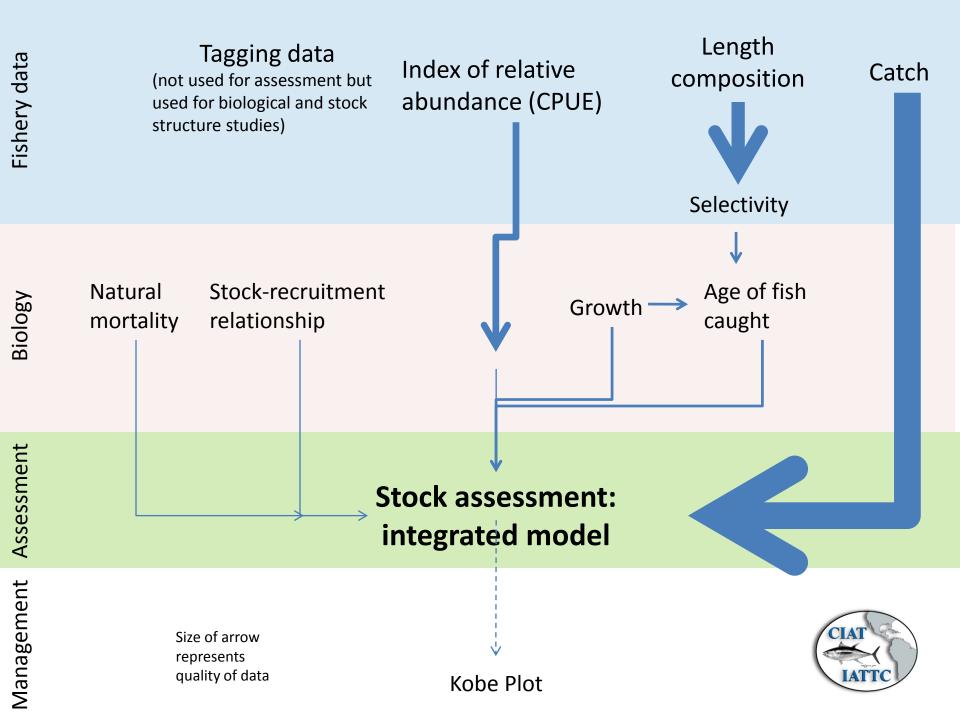
Integrated statistical age-structured population dynamics model

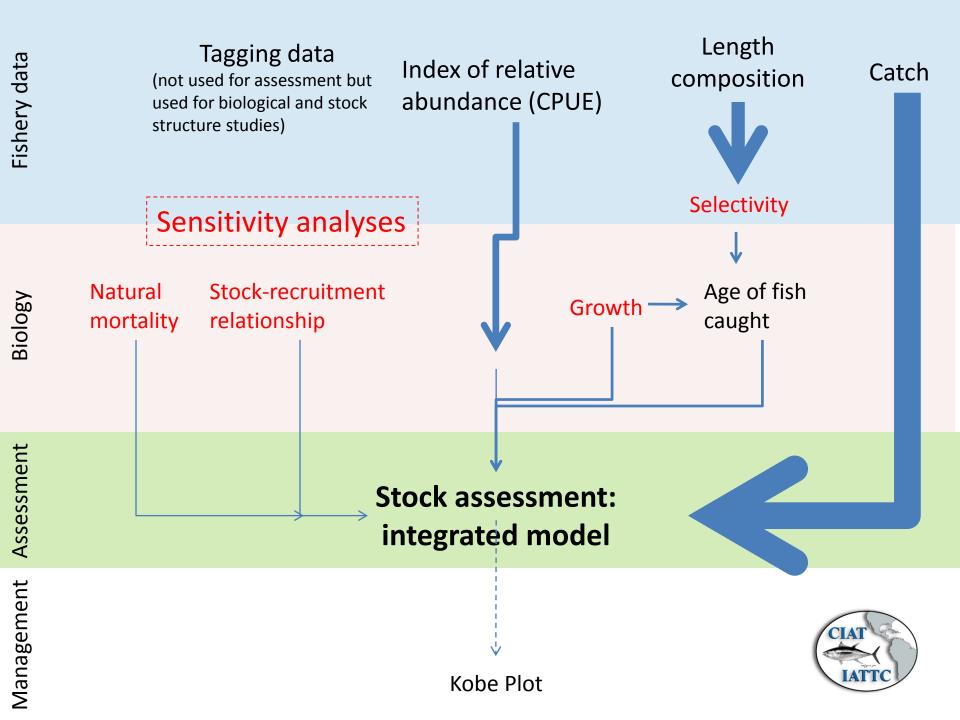
Assumptions:

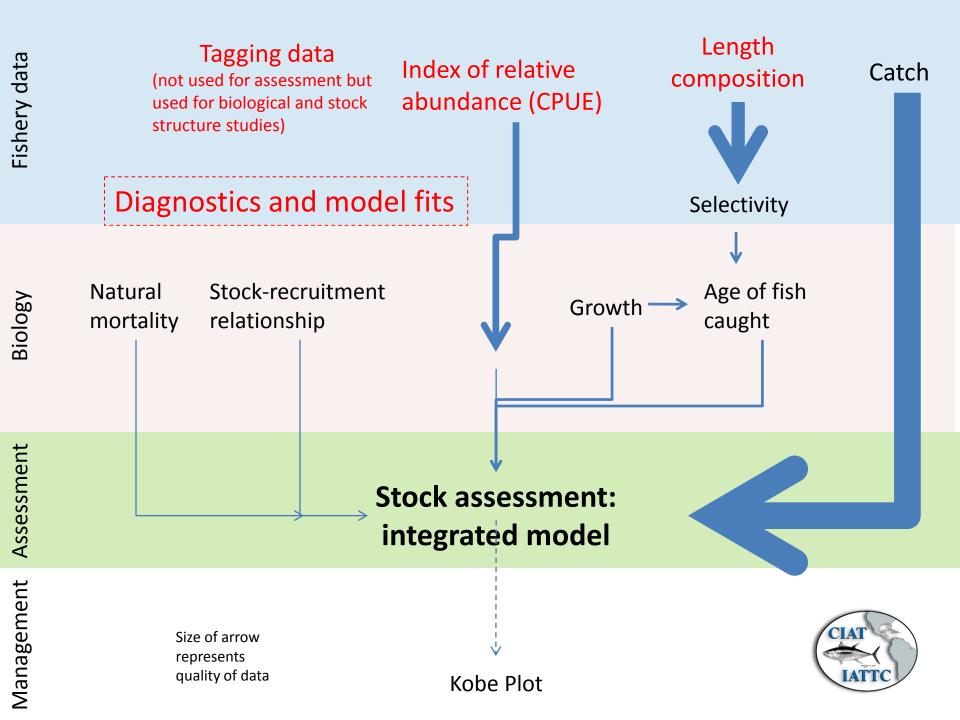
- one stock in the eastern Pacific Ocean
- 16 fisheries
- Sex-specific natural mortality
- One growth curve

• ...

Fit to indices of relative abundance and length-frequency data Using likelihood techniques





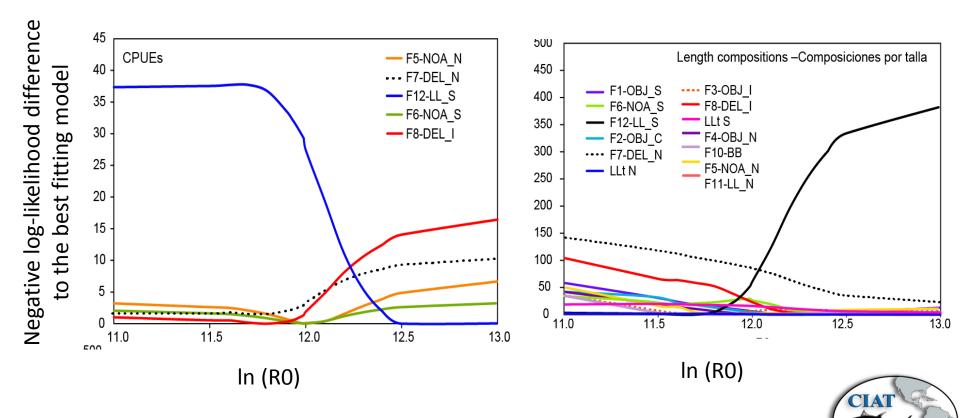


Diagnostics for integrated models

- Likelihood profile on the scaling parameter
- Age-structure production model fit only to indices of relative abundance
- Catch curve analysis fit only to length-frequency data
- Depletion model simplified population dynamics model, fit only to purse-seine CPUE

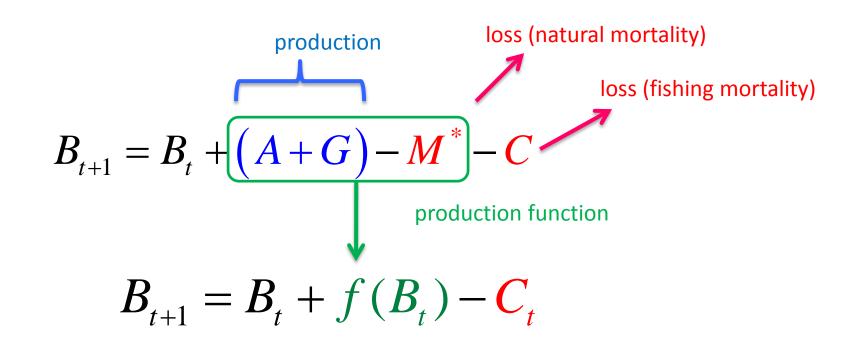
Diagnostics

RO is the virgin recruitment (the parameter that gives the absolute scale of the model)



Age-structure production model (ASPM)

Production and loss in a population



A production model aggregate the production and natural losses in a **production function**

- An age-structure production model takes into account how those processes change with **age** (size at age, natural mortality at age) and assumes a constant recruitment
- If the production function is constant, then the changes in biomass Bt are due to loss to fishing

 $Ct = f(Et \times Bt)$ Ct / Et = f(Bt)

- Ct / Et is the "catch per unit of effort" (CPUE)
- CPUE is proportional to the biomass of the stock Bt



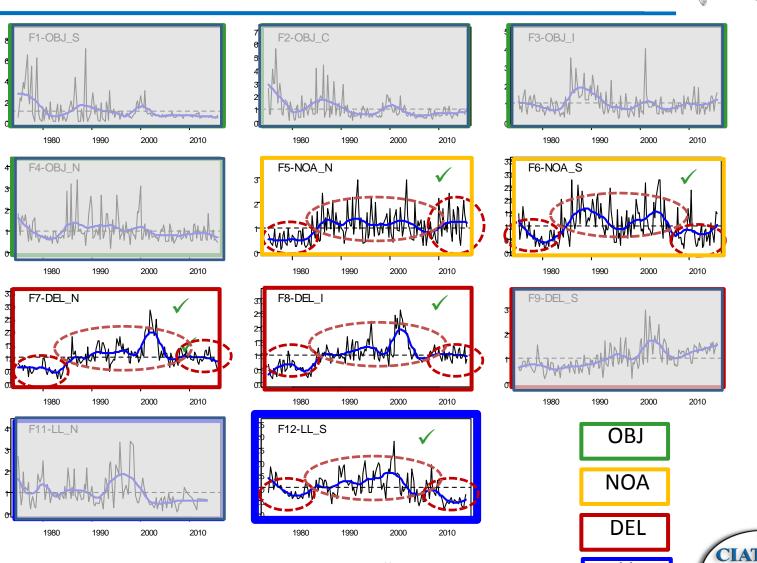
Fishery data

LL

IATTO

Fit to 5 CPUE series simultaneously

Scaled CPUE-CPUE escalade



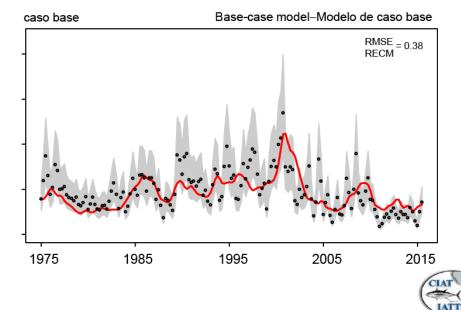
Year-año

Age-structured production model

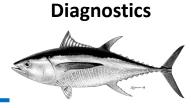


Diagnostics

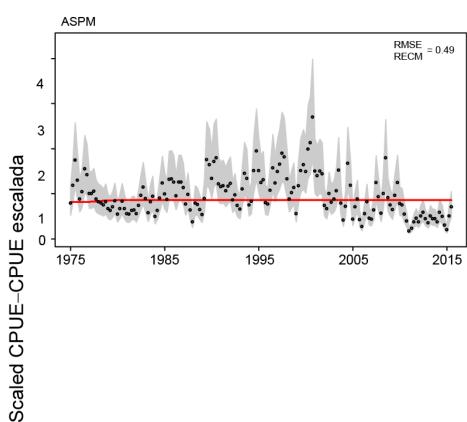
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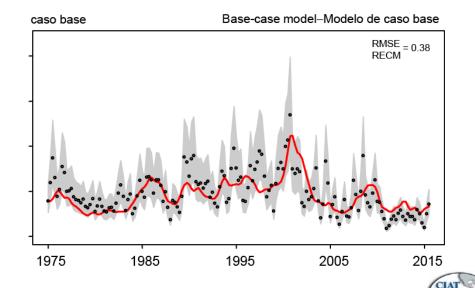


Age-structured production model

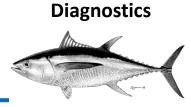


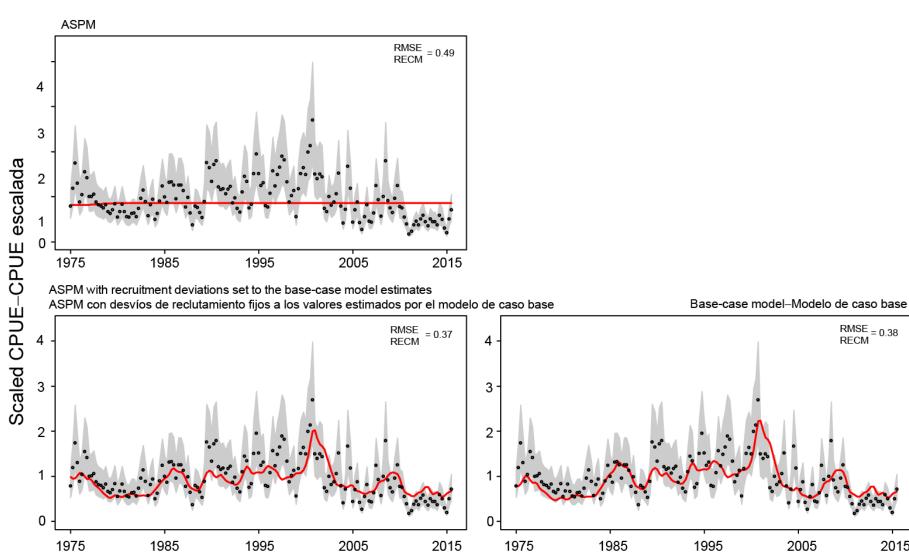
LATT





Age-structured production model



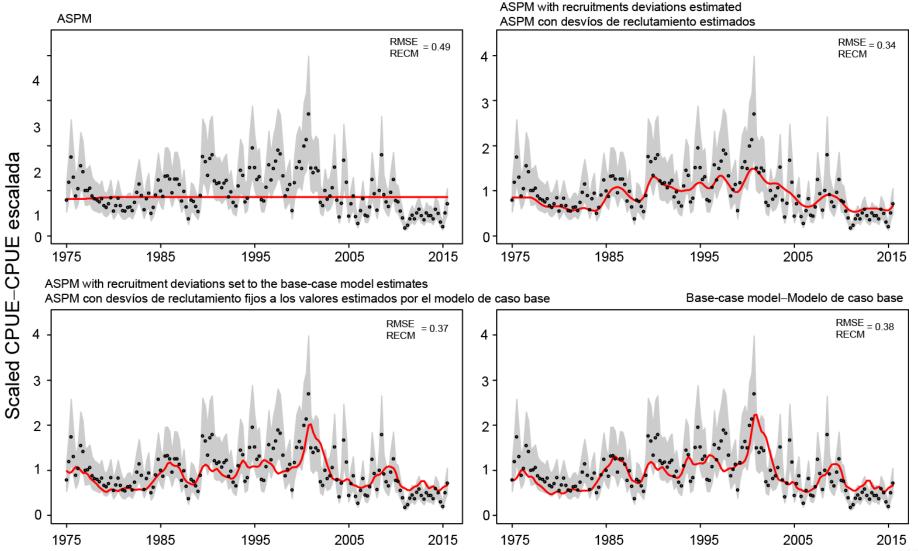




Diagnostics

Age-structured production model



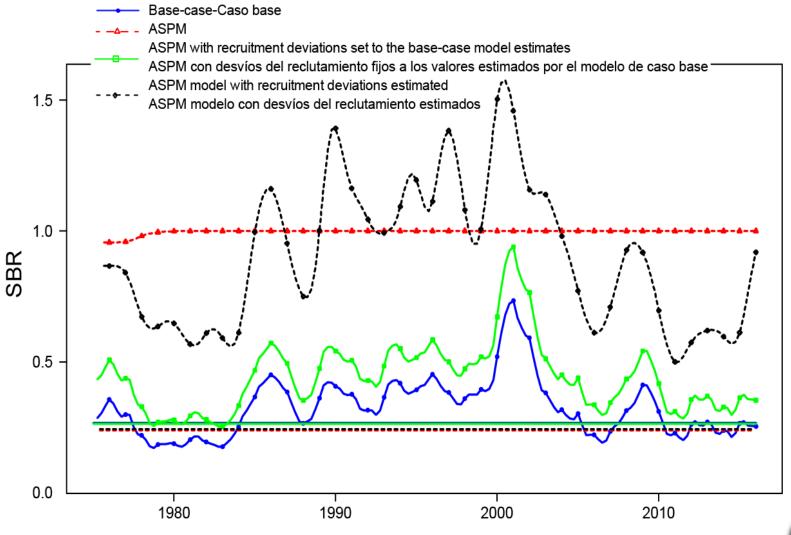




Diagnostics

Age-structured production model



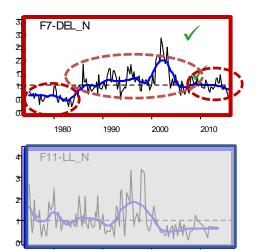




ASPM with one index only at a time



vJ

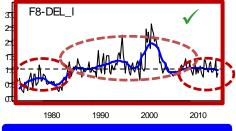


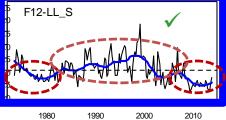
1990

2000

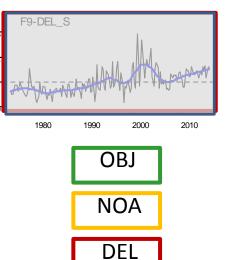
2010

1980

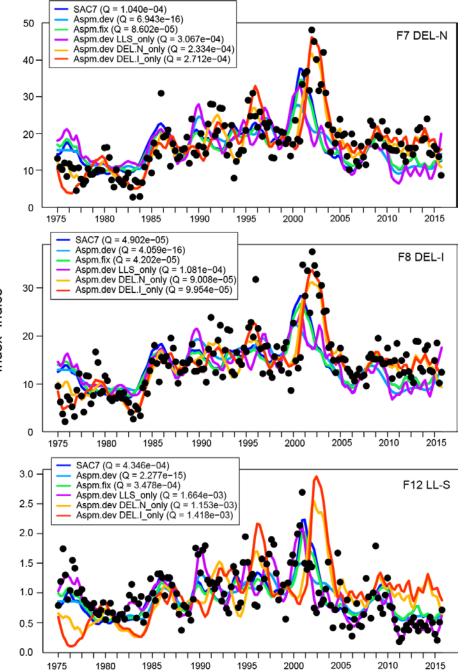




Year-año

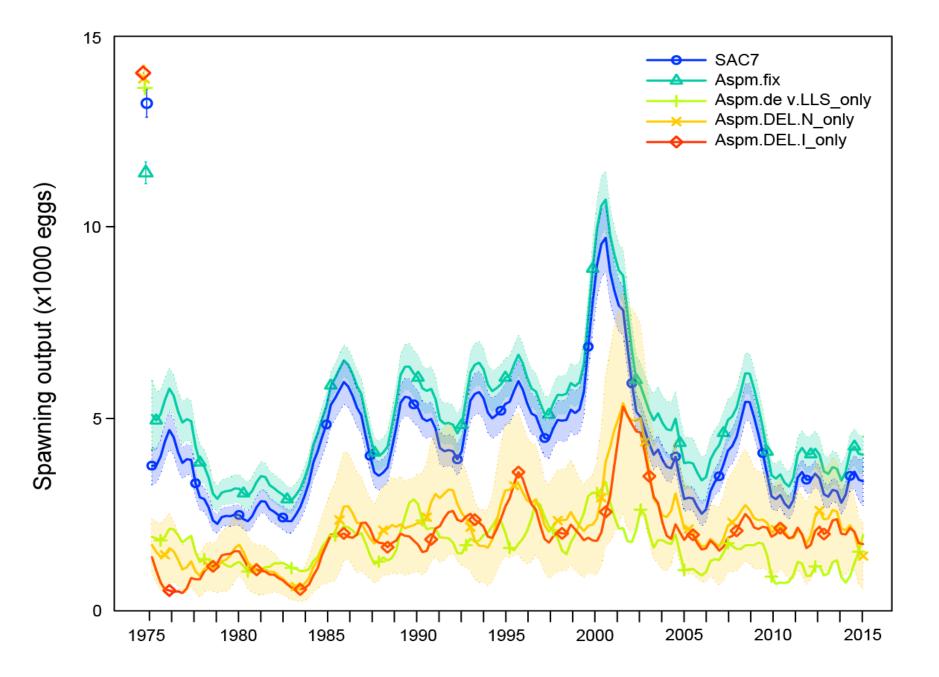






Fits to three CPUE series of the SAC7 base case model, the ASPMdev and the ASPMdev fit to one cpue series at a time

Index-Índice

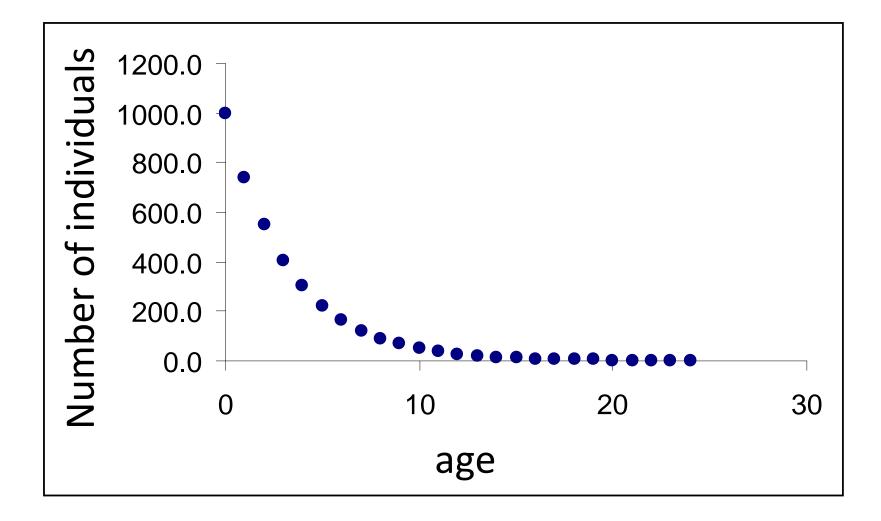


Results ASPM

- We cannot capture the biomass dynamics unless we take into account the recruitment deviations
 - However ASPMdev is only able to estimate the relative scale
 - Even when the recruitment estimates are provided, the model
 ASPMfix cannot estimate the same scale as the IM
- Fitting one index at a time allows to estimate a scale when recruitment deviations are estimated (ASPMdev)
 - The biomass is lower than what the IM predict
 - The uncertainty around estimates are high

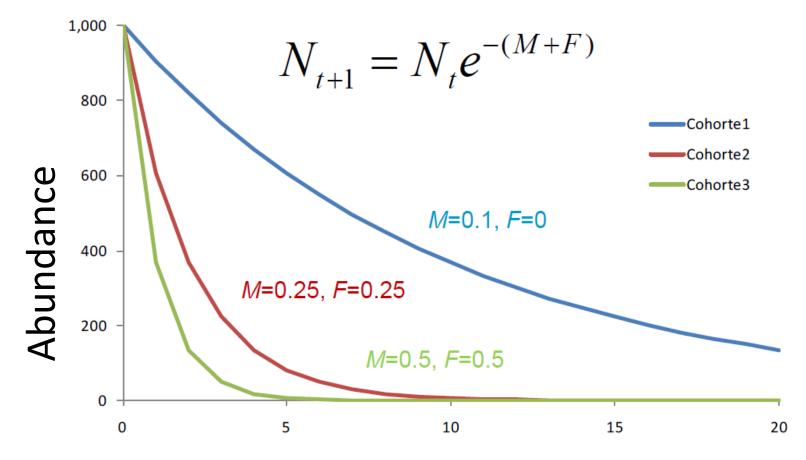
Catch-curve analysis

The trajectory of one cohort

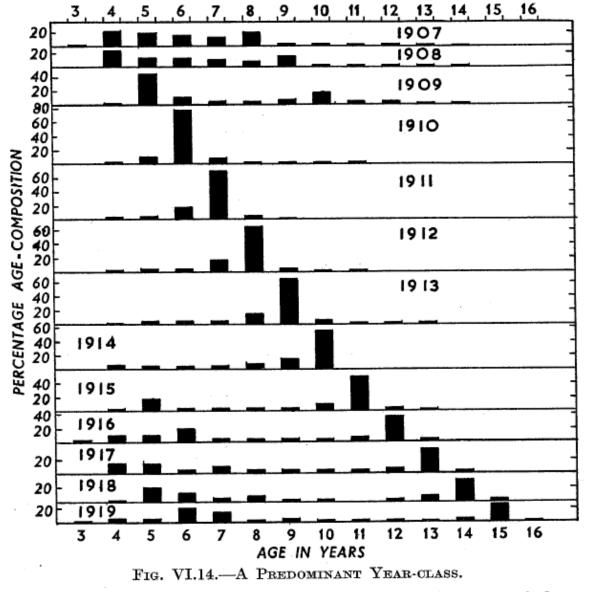


The trajectory of one cohort



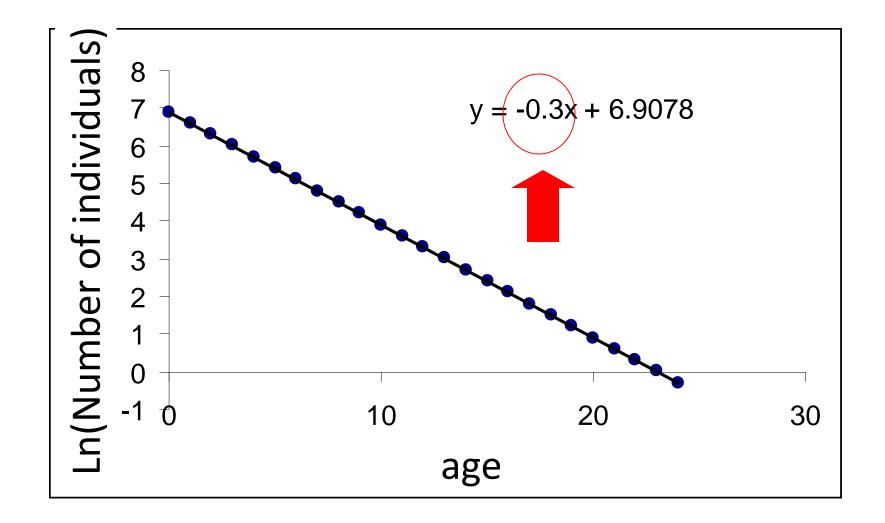


age

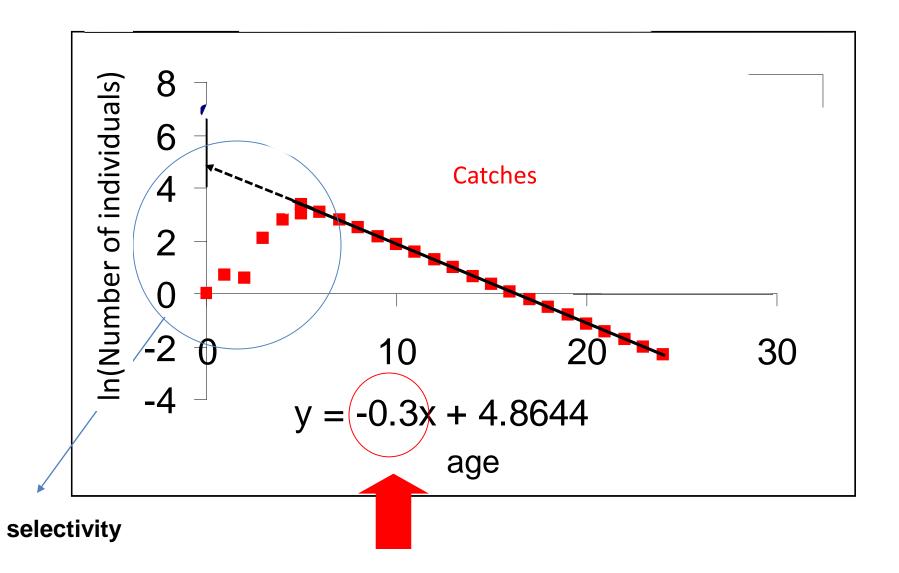


Percentage age composition of Norwegian herring in successive years of observation. (After Hjort, 1926)

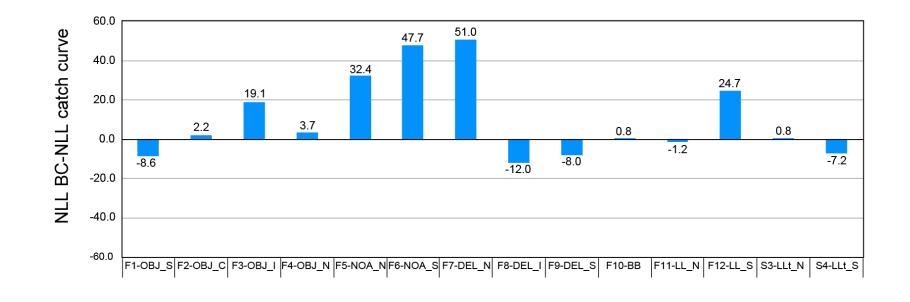
The trajectory of one cohort



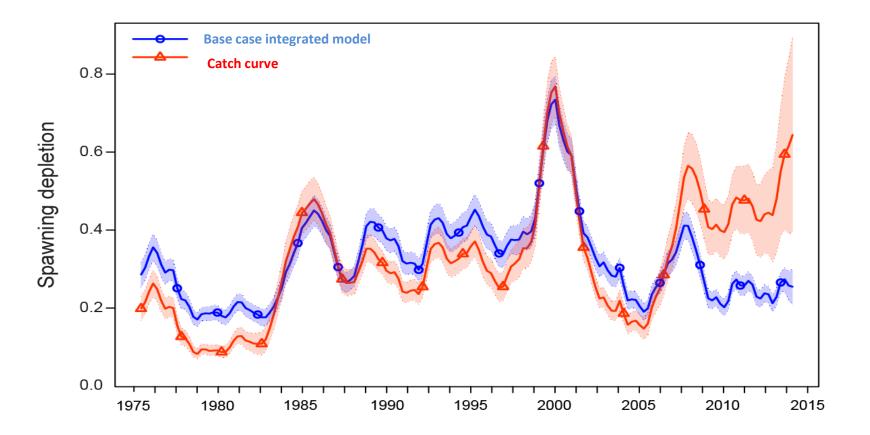
Catch curves



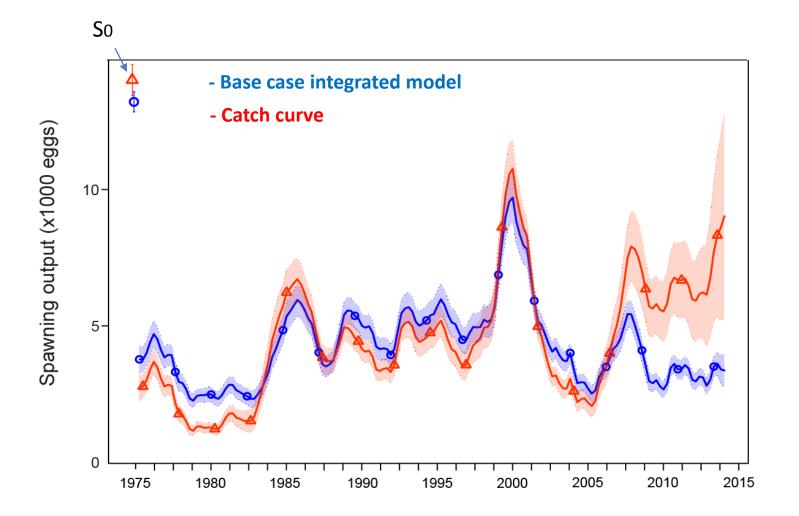
Results catch curve fits to length-frequency data

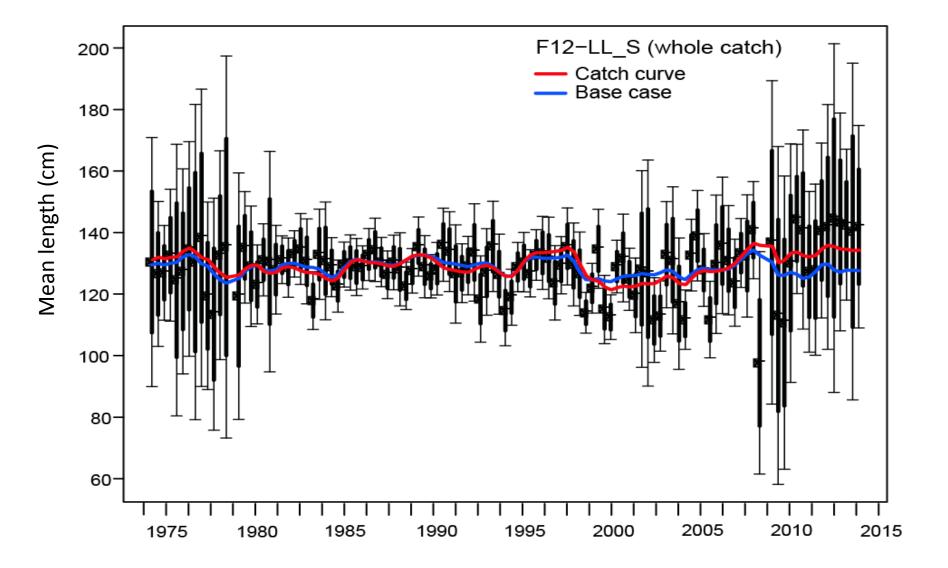


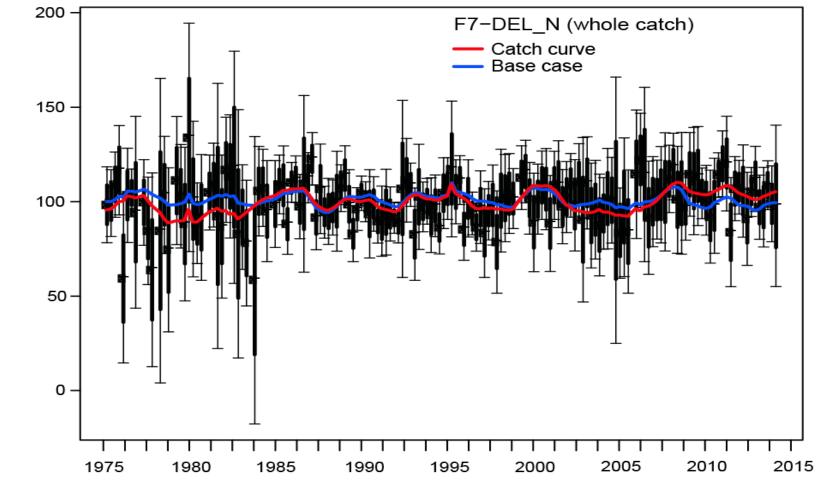
Results catch curve: relative abundance



Results catch curve: absolute abundance



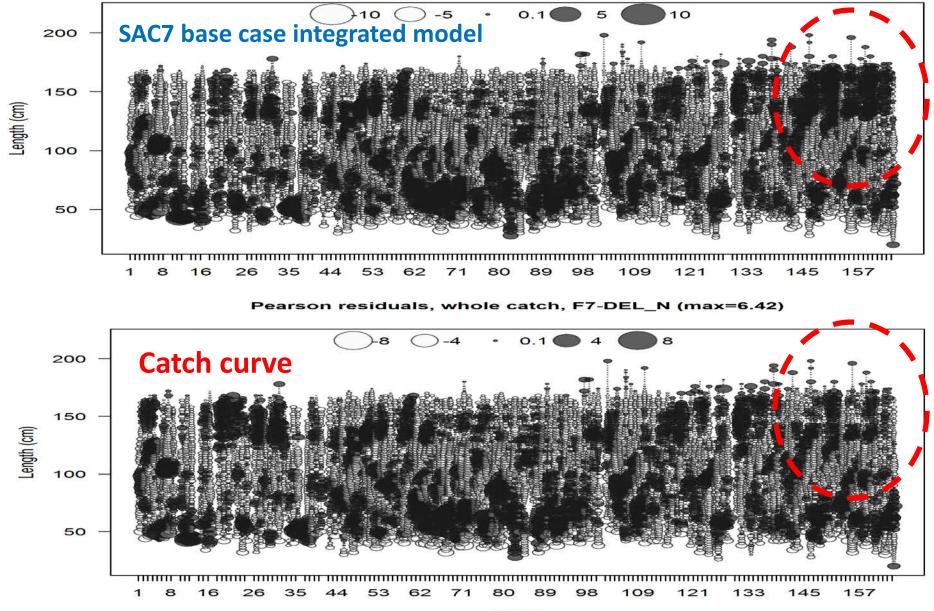




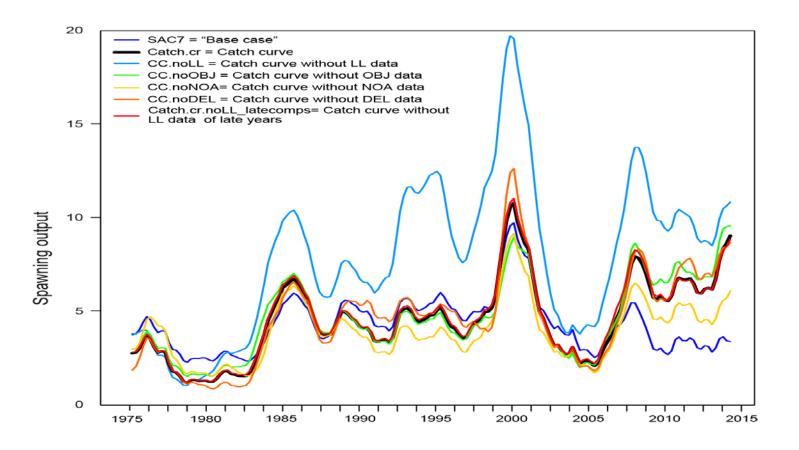
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Mean length (cm)

Pearson residuals, whole catch, F7-DEL_N (max=6.67)



Catch curve: sensitivity to data sets



Results catch curve

- The absolute biomass estimates from the catchcurve are similar to those of the base-case model for most of the series
- At the end of the series there is an indication model misspecification at the end of the time-series
- The length-frequency of LL_S fleet is in the most influent in the catch-curve results

Depletion model

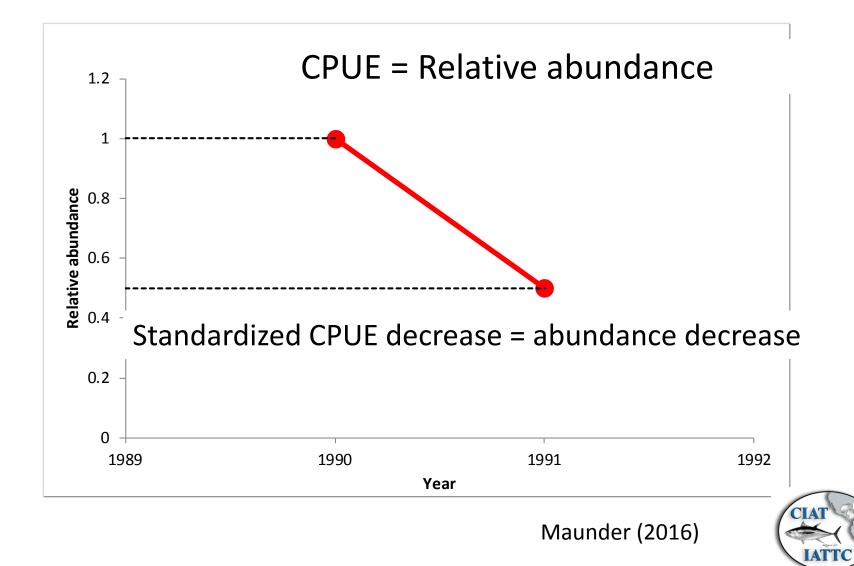
CPUE

 $Ct = q \times Et \times Bt$ $Ct / Et = q \times Bt$

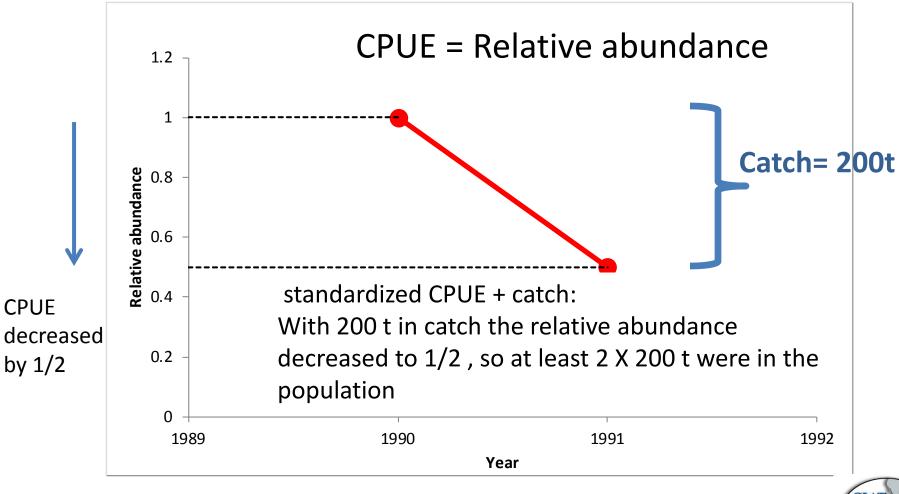
- Ct / Et is the "catch per unit of effort" (CPUE)
- CPUE is proportional to the biomass of the stock Bt
- The proportionality constant is **q** (catchability)



CPUE: why use it



CPUE: why use it



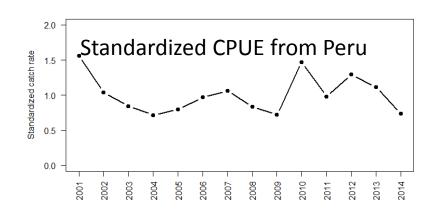
Maunder (2016)



CPUE: what scale? Dorado work

Annual cpue

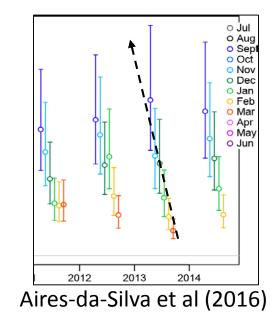
- Low information content
- Constant



Torrejón – Magallanes y Oliveros – Ramos (2015)

Monthly cpue

- High information content
- Fast growth
- High mortality rate



Standardized CPUE from Ecuador



Managing Data-Poor Fisheries: Case Studies, Models & Solutions 1:251–258, 2010 Copyright: California Sea Grant College Program 2010 ISBN number 978-1-888691-23-8

A Depletion Estimator for Within-Season Management of Yellowfin Tuna

MARK N. MAUNDER

Inter-American Tropical Tuna Commission, 8604 La Jolla Shores Drive, La Jolla, CA, 92037-1508, USA mmaunder@iattc.org

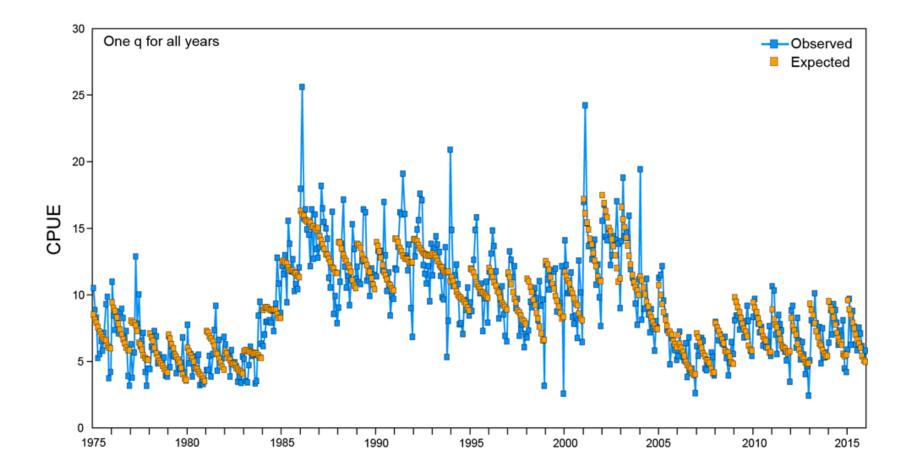
$$B_{y,w+1} = B_{y,w} (1+G)S + R - C_{y,w}$$

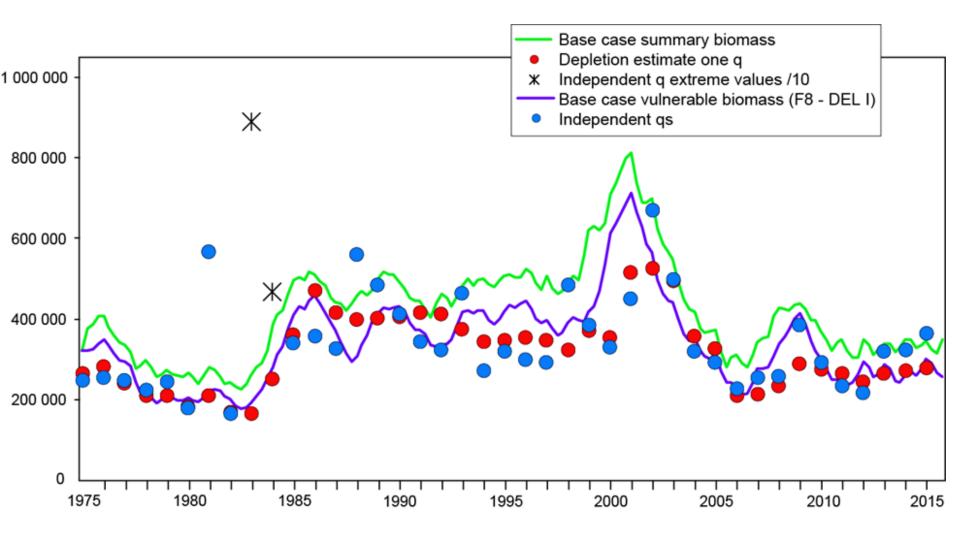
where $B_{y,w}$ is the biomass in week w of year y, and $C_{y,w}$ is the catch in week w of year y.

 $CPUE = q \times Bt$

In this study we used a monthly time step CPUE is purse-seine catch per day fished, regardless of set type

Results: fits to CPUE



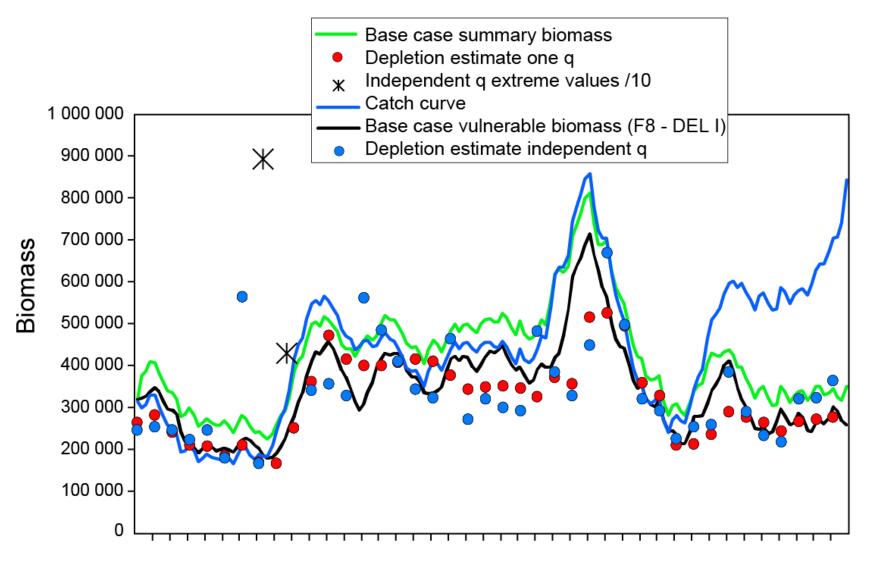


Results depletion model

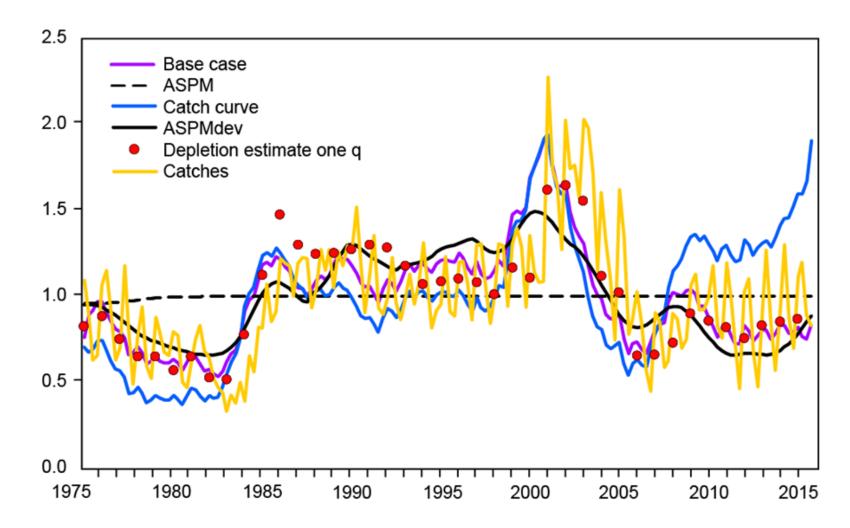
- The depletion estimator provided good fit to the data
- The depletion model was able to provide an estimate of the absolute abundance, which coincided with the base-case model, this increased our confidence on the integrated model
- This approach should investigated for improving the assessment, for example by changing to a monthly time step

Summary of results

Absolute abundance



Relative abundance

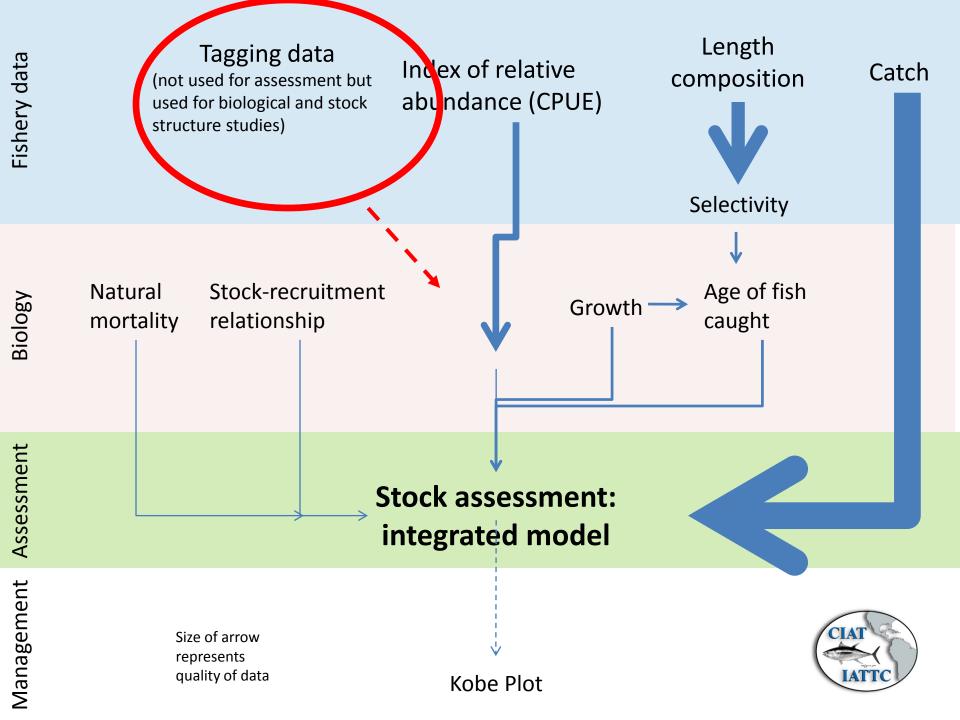


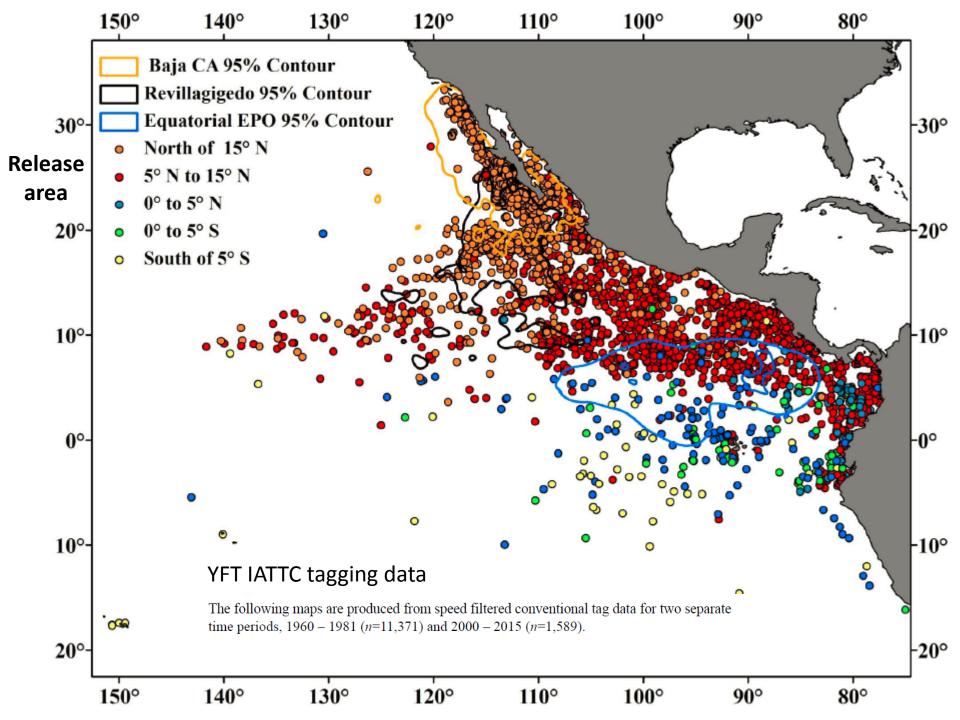
Conclusions

- The absolute scale cannot be estimated from the five indices of relative abundance (CPUE) alone within the age-structure production model, unless precise estimates of recruitment deviations are provided (ASPMfix)
- Only relative changes in abundance can be estimated from the five indices of relative abundance (CPUE) within a ASPMdev model
- The indices of relative abundance from LL are in contradiction to those from PS
- The LL index should be investigated for changes in the fishery that might have influenced its ability to represent relative abundance.

Conclusions

- The absolute biomass estimates from the catch-curve are similar to those of the base-case model, except at the end of the series, which might be an indication model misspecification at the end of the time-series
- The depletion estimator provided good fit to the data, was able to provide an estimate of the absolute abundance, which coincided with the base-case model, and increase our confidence in the stock assessment model.
- Given the good performance of the depletion model presented here and in Maunder (2010), suggests that the approach should investigated for improving the assessment, for example by changing to a monthly time step
- Also, the depletion model could provide "in-season" abundance estimates, important if catch quotas are used for management





Conclusions

- Tagging data could be another piece of information to be integrated into the models
- Tagging data provides information on biological (growth, natural mortality, movement and mixing rates) and fisheries process
- A large-scale tagging program can be used to estimate abundance

Thank you