# How to insert r4ss Figure

Melissa Monk

#### Data by type and year

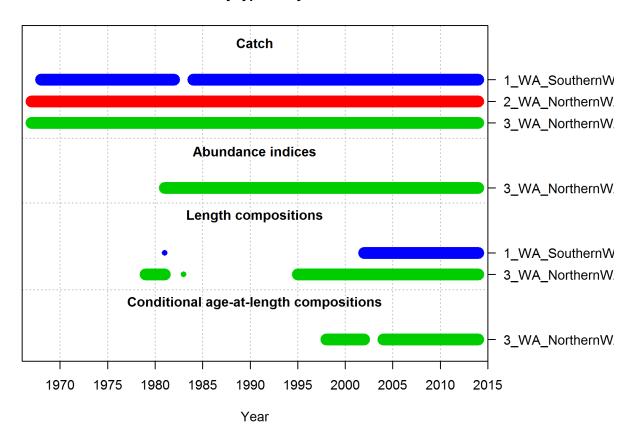


Figure 1: Summary of data sources used in the base case assessment. fig:data\_plot

### Ending year expected growth (with 95% intervals)

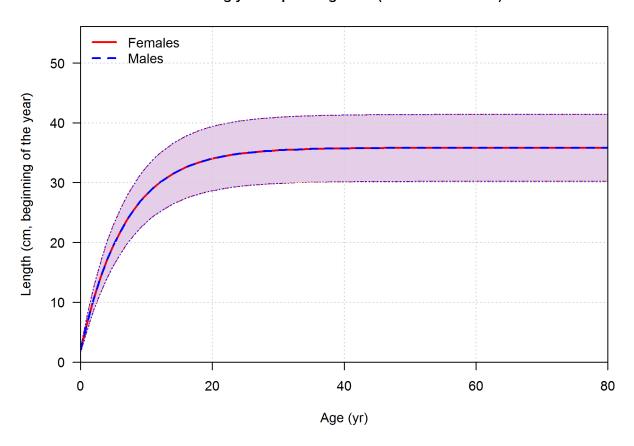


Figure 2: Length at age in the beginning of the year (or season) in the ending year of the model. Shaded area indicates 95% distribution of length at age around estimated growth curve. fig:modi\_1\_bio1\_sizeatage

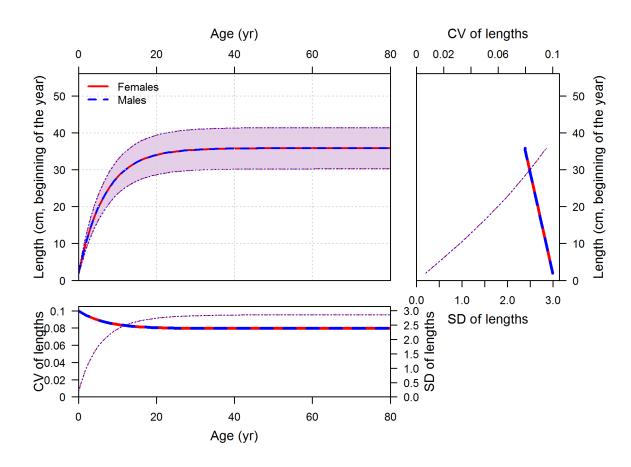


Figure 3: Length at age (top\_left panel) with CV (thick line) and SD (thin line) of length at age shown in top\_right and lower\_left panels | fig:mod1\_2\_bio2\_sizeatage\_plus\_CV\_and\_SD |

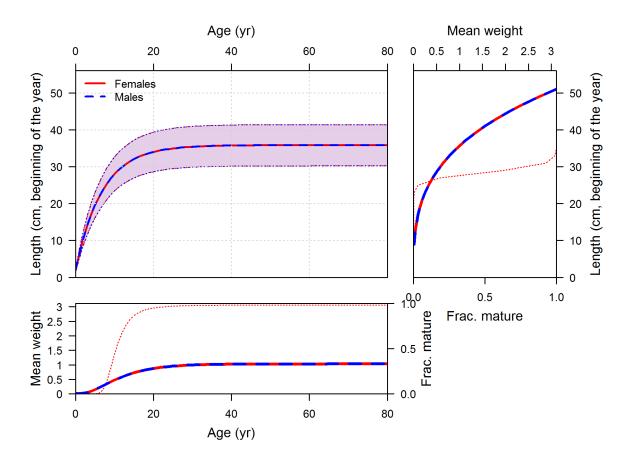


Figure 4: Length at age (top\_left panel) with weight (thick line) and maturity (thin line) shown in top\_right and lower\_left panels | fig:mod1\_3\_bio3\_sizeatage\_plus\_WT\_and\_MAT

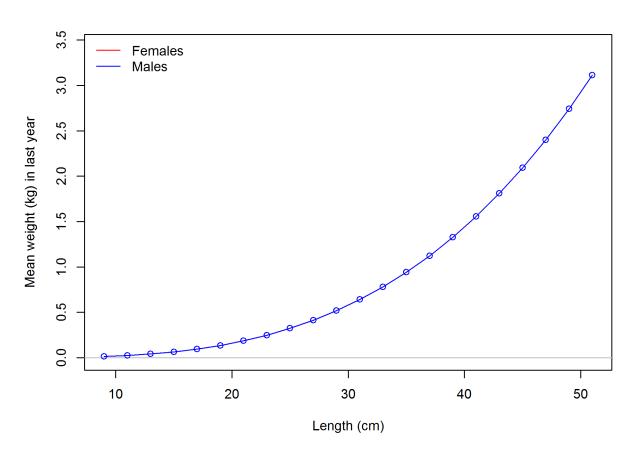


Figure 5: Weight\_length relationship for females  $^{\texttt{fig:mod1\_4\_bio4\_weightatsize}}$ 

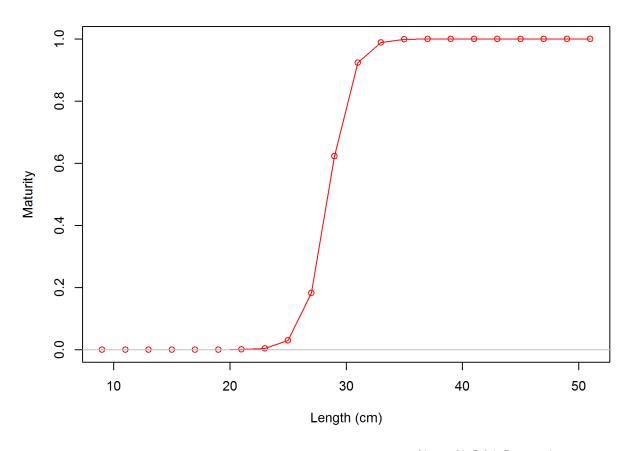


Figure 6: Maturity at length  $fig:mod1_5_bio5_maturity$ 

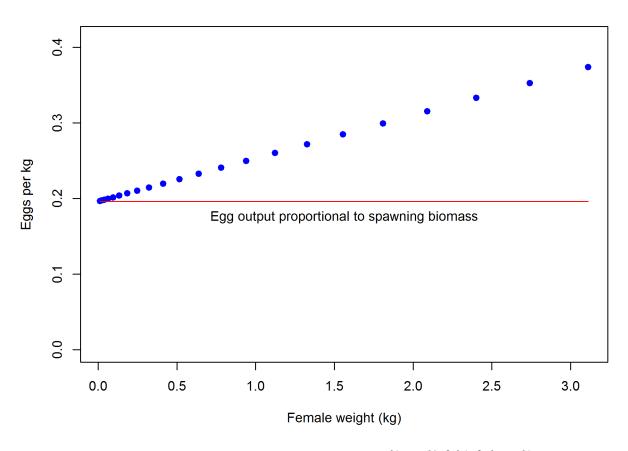


Figure 7: Fecundity fig:mod1\_6\_bio6\_fecundity

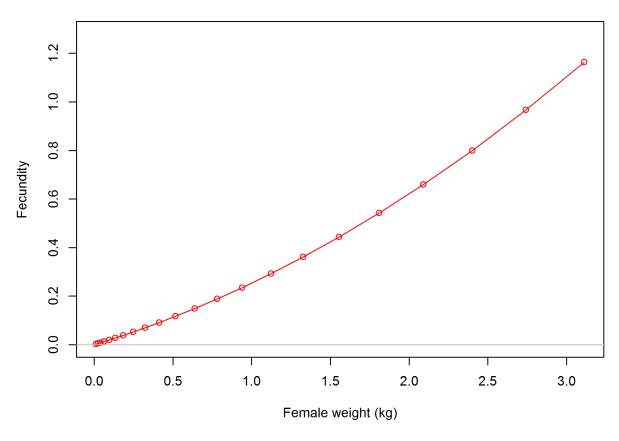


Figure 8: Fecundity as a function of weight  $\lceil \text{ig:mod1\_7\_bio7\_fecundity\_wt} \rceil$ 

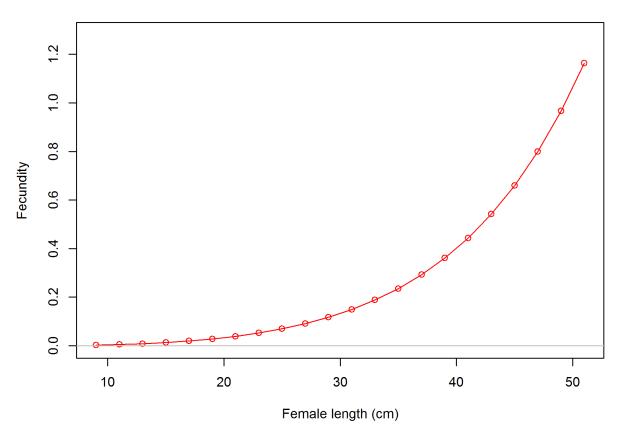


Figure 9: Fecundity as a function of length  $\lceil \text{fig:mod1\_8\_bio8\_fecundity\_len} \rceil$ 

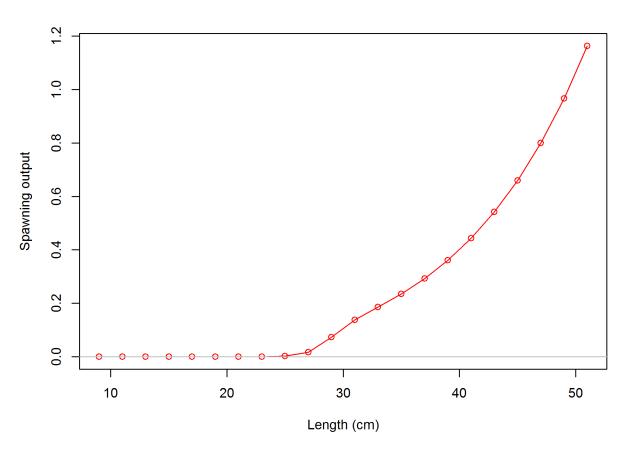
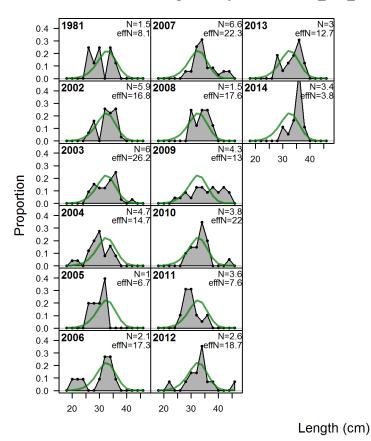


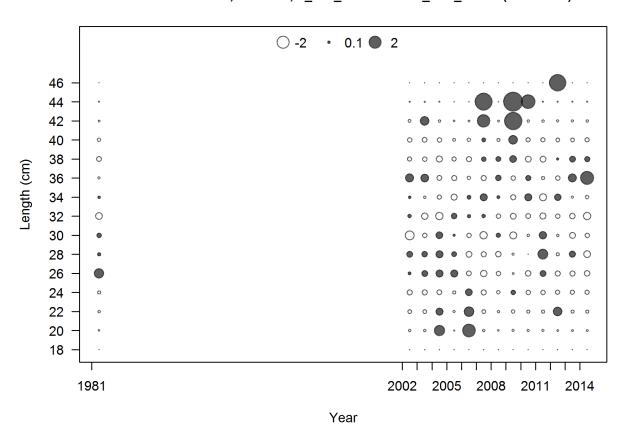
Figure 10: Spawning output at length | fig:mod1\_9\_bio9\_spawningoutput

### length comps, retained, 1\_WA\_SouthernWA\_Rec\_PCPR

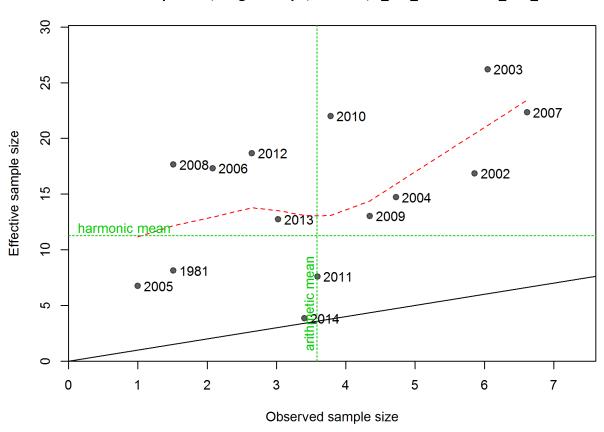


 $Figure~11:~length~comps,~retained,~1\_WA\_SouthernWA\_Rec\_PCPR~ \\ \\ ^{\texttt{fig:mod1\_1\_comp\_lenfit\_flt1mkt2}}$ 

### Pearson residuals, retained, 1\_WA\_SouthernWA\_Rec\_PCPR (max=4.76)



### N-EffN comparison, length comps, retained, 1\_WA\_SouthernWA\_Rec\_PCPR



 $Figure~13:~N\_EffN~comparison,~length~comps,~retained,~1\_WA\_SouthernWA\_Rec\_PCPR~\\ \\ [fig:mod1\_3\_comp\_lenfit] \\ [fig:mod1\_3\_comp$ 

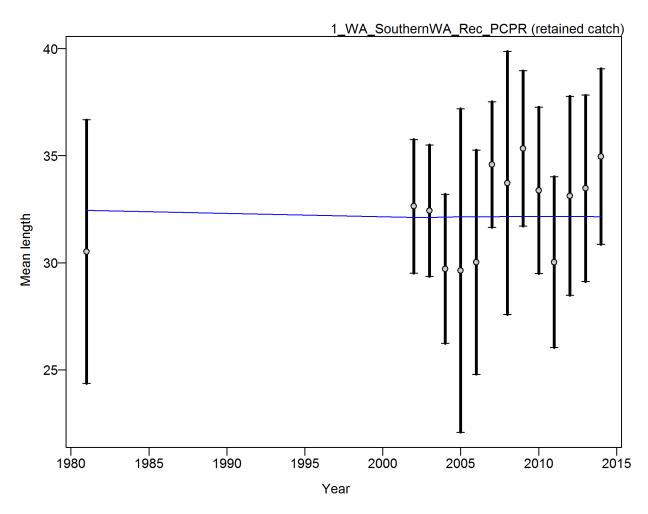
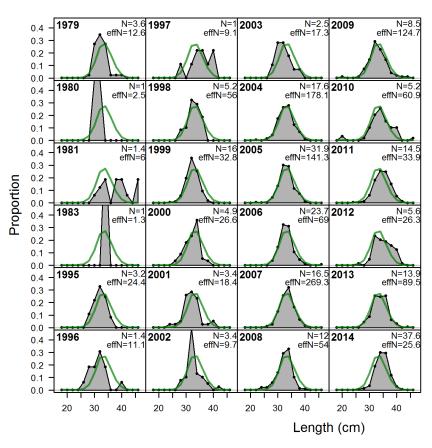


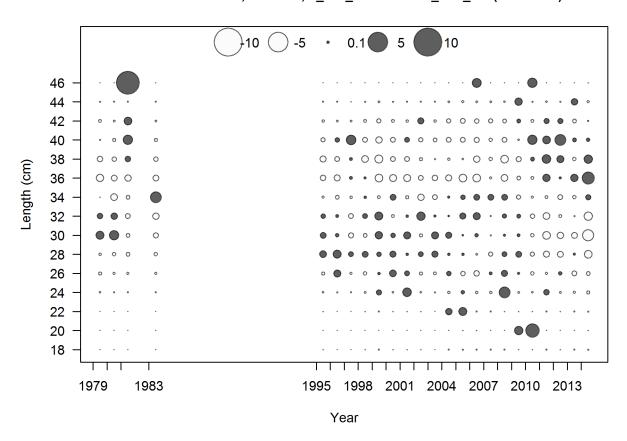
Figure 14: Francis data weighting method TA1.8 1\_WA\_SouthernWA\_Rec\_PCPR Suggested sample size adjustment (with 95% interval) for len data from 1\_WA\_SouthernWA\_Rec\_PCPR: 0.9991 (0.6951\_2.1587) fig:mod1\_4\_comp\_lenfit\_data\_weighting\_TA1.8\_1\_WA\_SouthernWA\_Rec\_PCPR

### length comps, retained, 3\_WA\_NorthernWA\_Rec\_PR

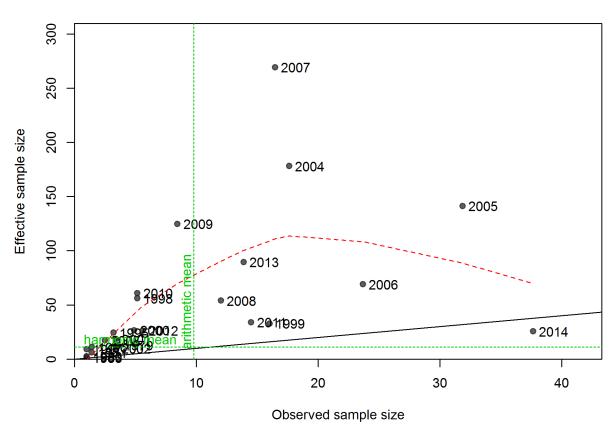


 $Figure~15:~length~comps,~retained,~3\_WA\_NorthernWA\_Rec\_PR \\ ~ fig:mod1\_5\_comp\_lenfit\_flt3mkt2 \\$ 

### Pearson residuals, retained, 3\_WA\_NorthernWA\_Rec\_PR (max=6.82)



# N-EffN comparison, length comps, retained, 3\_WA\_NorthernWA\_Rec\_PR



 $Figure~17:~N\_EffN~comparison,~length~comps,~retained,~3\_WA\_NorthernWA\_Rec\_PR~ \\ fig:mod1\_7\_comp\_lenfit\_started,~2\_WA\_NorthernWA\_Rec\_PR~ \\ fig:mod1\_7\_comp\_lenfit\_started,~2\_WA\_NORTHernWA\_R$ 

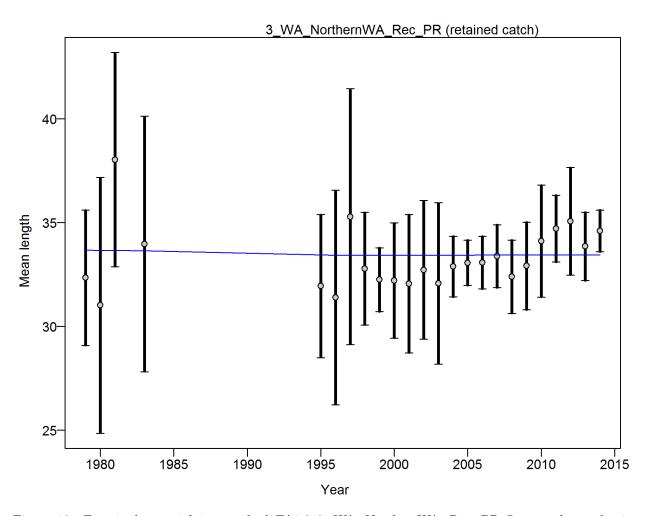
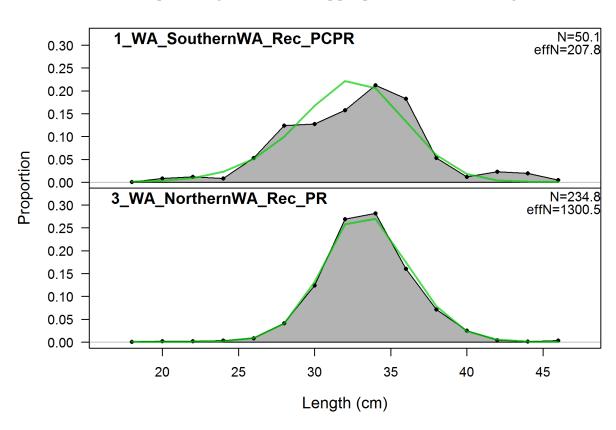


Figure 18: Francis data weighting method TA1.8 3\_WA\_NorthernWA\_Rec\_PR Suggested sample size adjustment (with 95% interval) for len data from 3\_WA\_NorthernWA\_Rec\_PR: 0.9797 (0.6452\_2.3201) fig:mod1\_8\_comp\_displayed adjustment (with 95% interval) for len data from 3\_WA\_NorthernWA\_Rec\_PR: 0.9797 (0.6452\_2.3201)

# length comps, retained, aggregated across time by fleet



 $\label{eq:fig:mod1_9_comp_lenfit_mkt2_aggregated} Figure \ 19: \ length \ comps, \ retained, \ aggregated \ across \ time \ by \ fleet \ {\tt fig:mod1_9\_comp\_lenfit\_mkt2\_aggregated}$ 

### Pearson residuals, sexes combined, retained, comparing across fleets

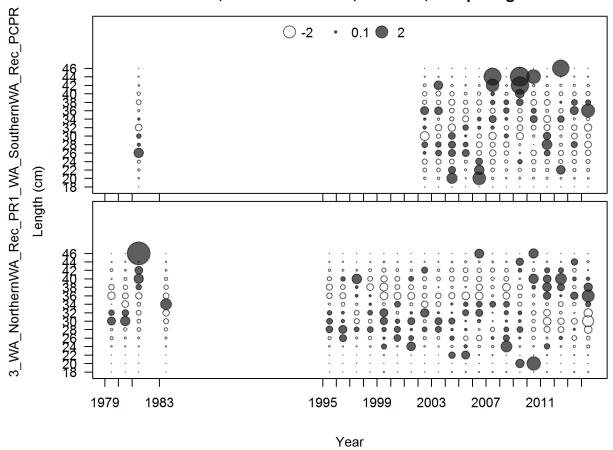
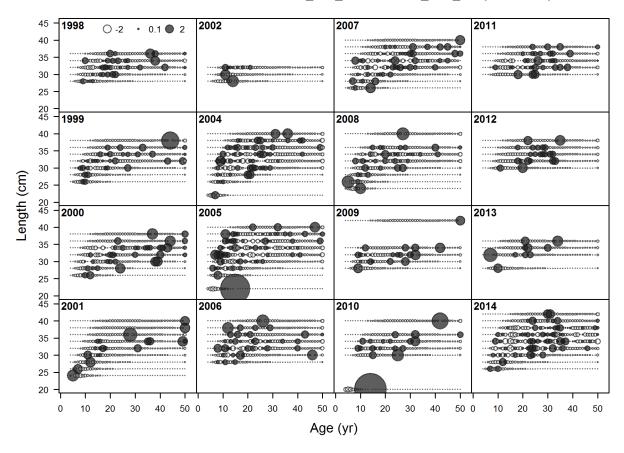


Figure 20: Note: this plot doesn't seem to be working right for some models. Pearson residuals, sexes combined, retained, comparing across fleets Closed bubbles are positive residuals (observed > expected) and open bubbles are negative residuals (observed < expected). fig:mod1\_10\_comp\_lenfit\_sex1mkt2\_multi-fleet\_comparison

# Pearson residuals, retained, 3\_WA\_NorthernWA\_Rec\_PR (max=29.9)



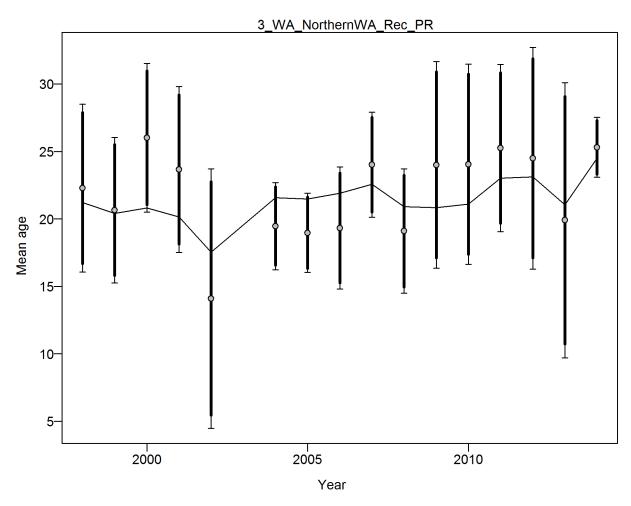


Figure 22: Francis data weighting method TA1.8 for conditional age data 3\_WA\_NorthernWA\_Rec\_PR For more info, see Francis, R.I.C.C. (2011). Data weighting in statistical fisheries stock assessment models. Can. J. Fish. Aquat. Sci. 68: 1124\_1138. [fig:mod1\_2\_comp\_condAlfit\_data\_weighting\_TA1.8\_condAge3\_WA\_NorthernWA\_Rec\_Can. J. Fish. Aquat. Sci. 68: 1124\_1138.]

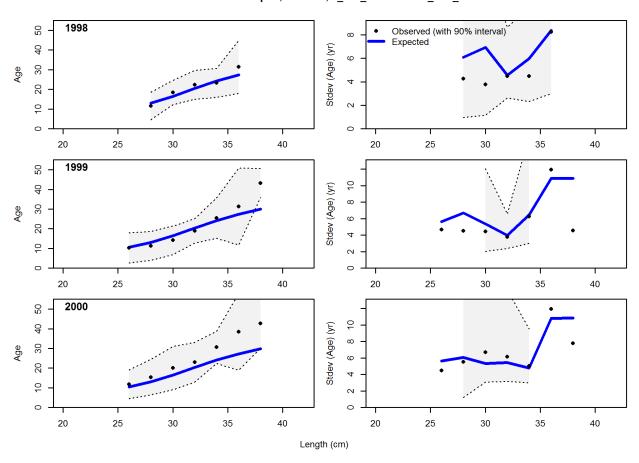


Figure 23: Conditional AAL plot, retained, 3\_WA\_NorthernWA\_Rec\_PR (plot 1 of 6) These plots show mean age and std. dev. in conditional AAL. Left plots are mean AAL by size\_class (obs. and pred.) with 90% CIs based on adding 1.64 SE of mean to the data. Right plots in each pair are SE of mean AAL (obs. and pred.) with 90% CIs based on the chi\_square distribution. fig:mod1\_3\_comp\_condAALfit\_Andre\_plotsflt3mkt2\_page1

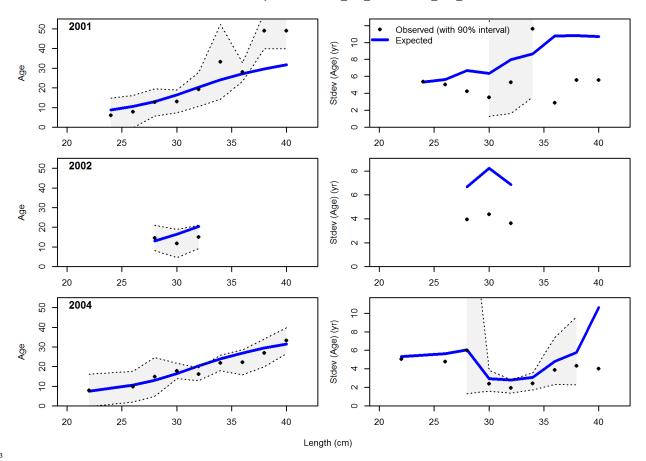


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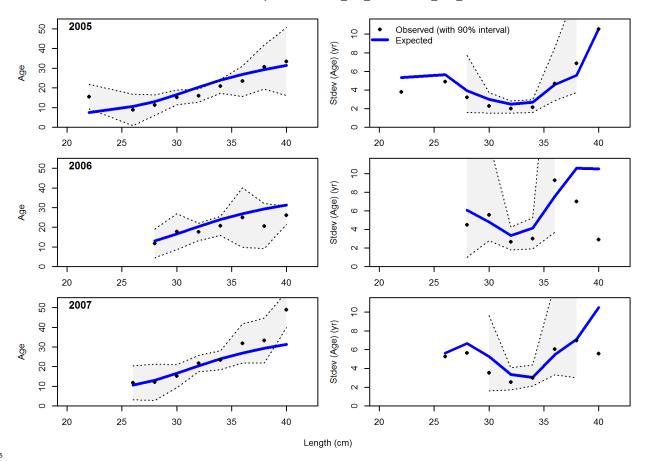


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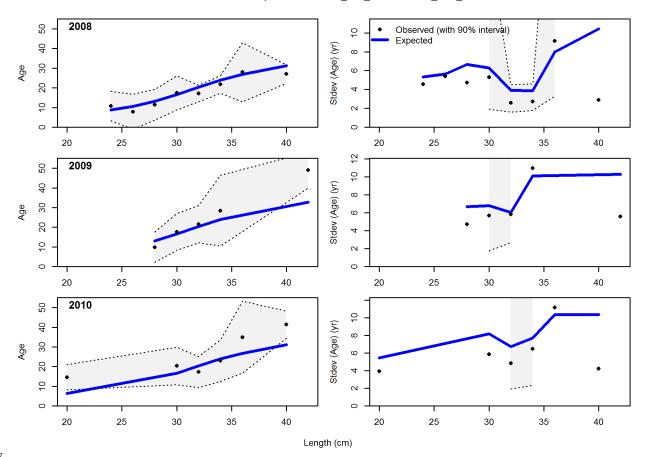


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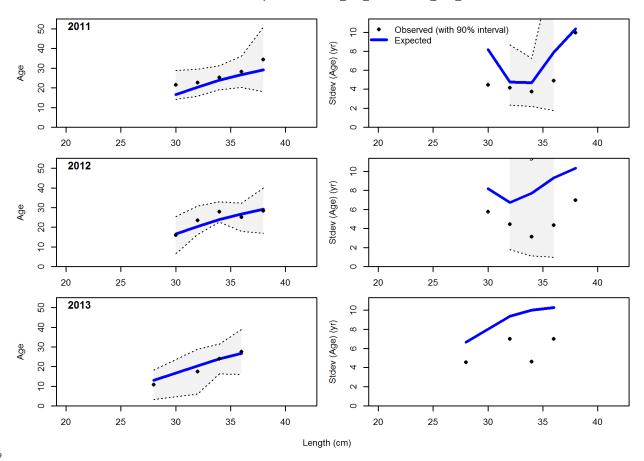
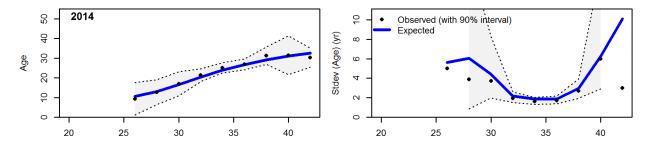


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Length (cm)

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12

### Data by type and year

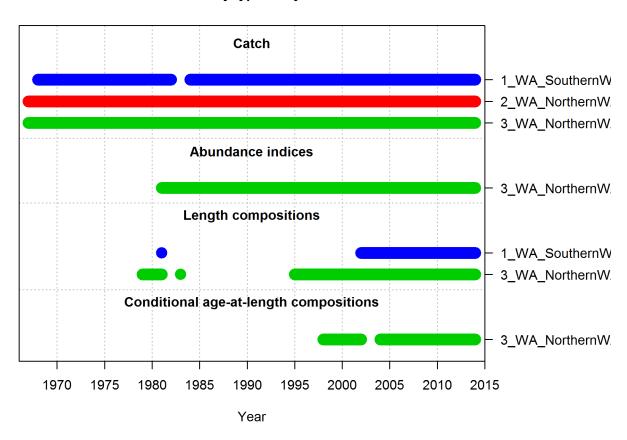


Figure 24: 1-Summary of data sources used in the base case assessment. fig:data\_plot

### Data by type and year

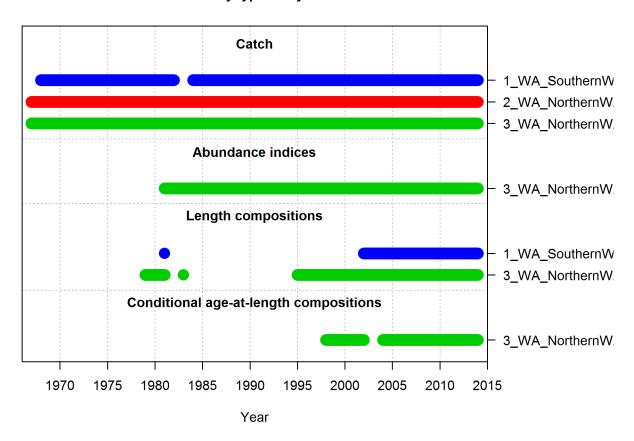


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# Data by type and year

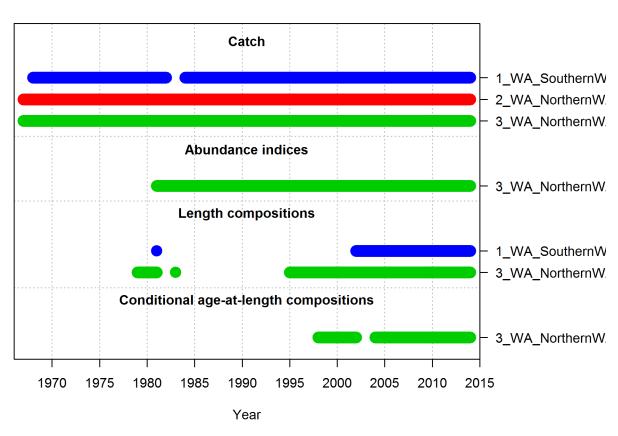


Figure 25: 3-Summary of data sources used in the base case assessment. fig:data\_plot