r4ss Figure Examples

Melissa Monk July 26, 2016

Data by type and year

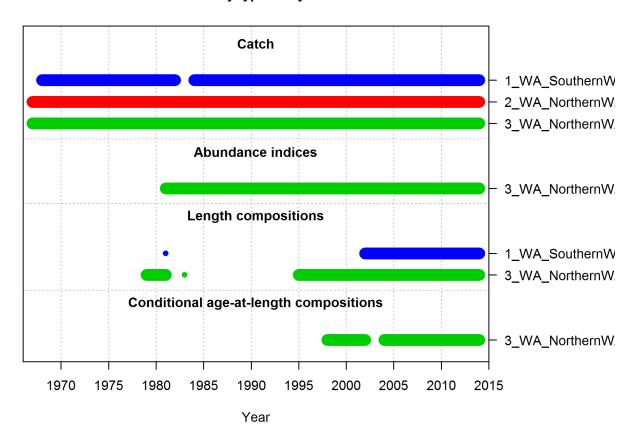


Figure 1: Summary of data sources used in the base case assessment. fig:data_plot

Ending year expected growth (with 95% intervals)

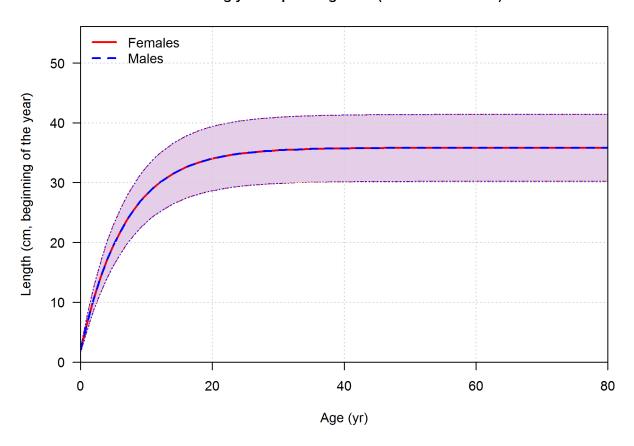


Figure 2: Length at age in the beginning of the year (or season) in the ending year of the model. Shaded area indicates 95% distribution of length at age around estimated growth curve. fig:modi_1_bio1_sizeatage

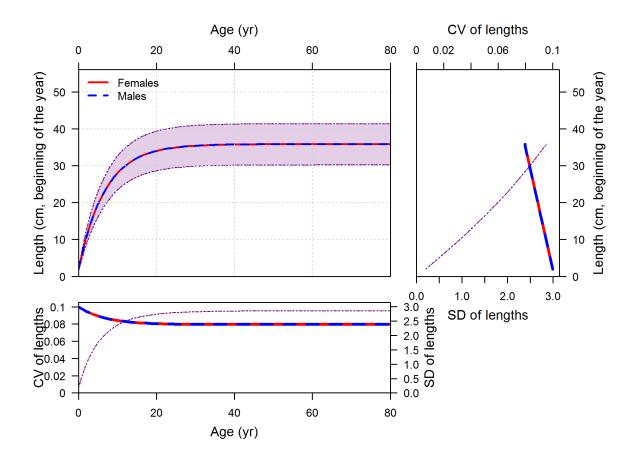


Figure 3: Length at age (top_left panel) with CV (thick line) and SD (thin line) of length at age shown in top_right and lower_left panels | fig:mod1_2_bio2_sizeatage_plus_CV_and_SD |

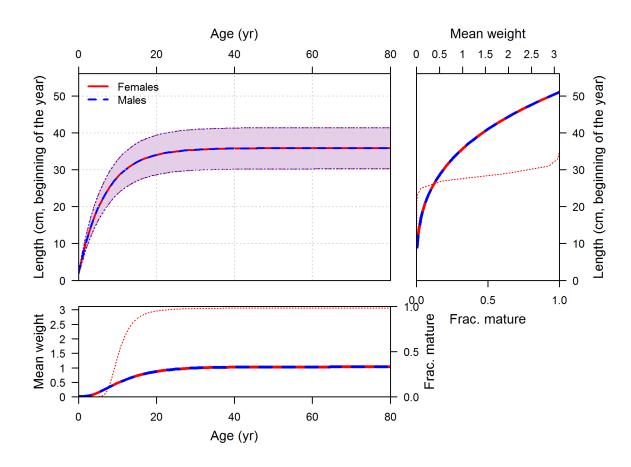


Figure 4: Length at age (top_left panel) with weight (thick line) and maturity (thin line) shown in top_right and lower_left panels | fig:mod1_3_bio3_sizeatage_plus_WT_and_MAT

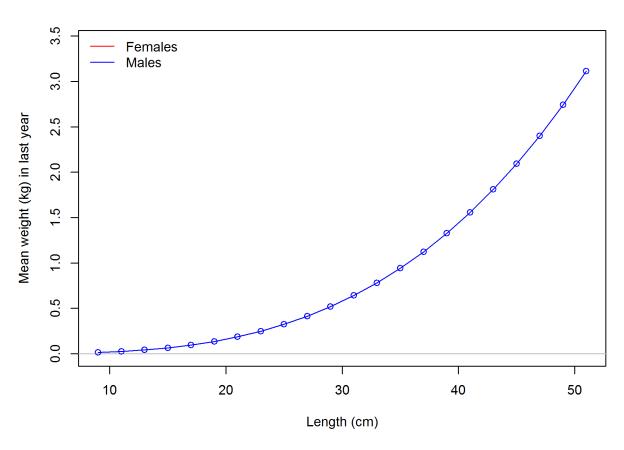


Figure 5: Weight_length relationship for females $^{\texttt{fig:mod1_4_bio4_weightatsize}}$

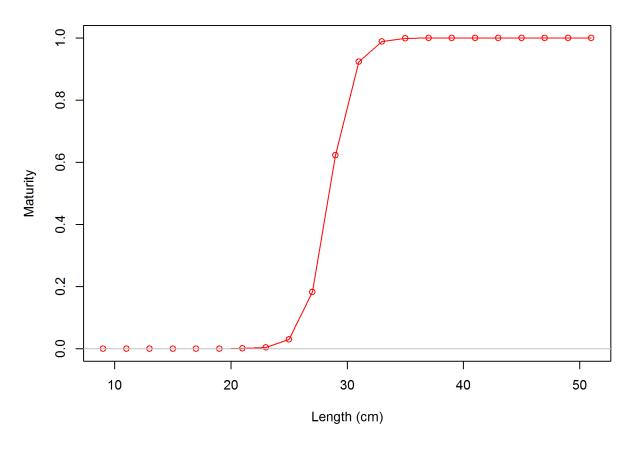


Figure 6: Maturity at length $fig:mod1_5_bio5_maturity$

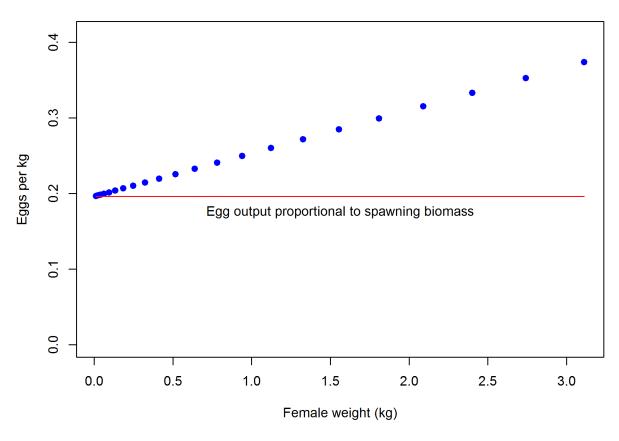


Figure 7: Fecundity fig:mod1_6_bio6_fecundity

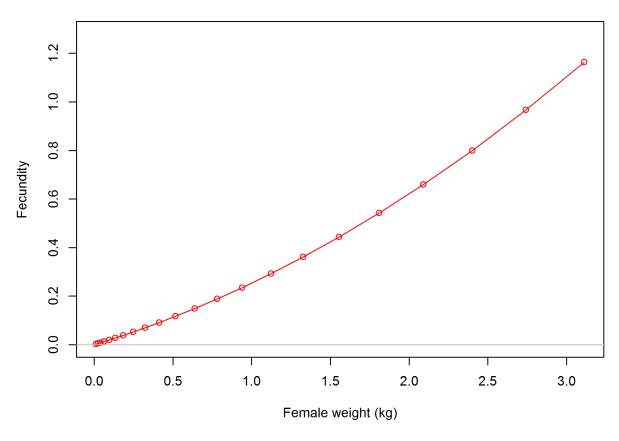


Figure 8: Fecundity as a function of weight $\lceil \text{ig:mod1_7_bio7_fecundity_wt} \rceil$

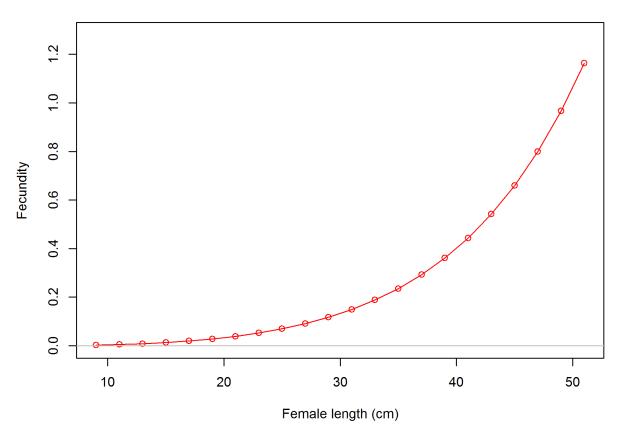


Figure 9: Fecundity as a function of length $\lceil \text{fig:mod1_8_bio8_fecundity_len} \rceil$

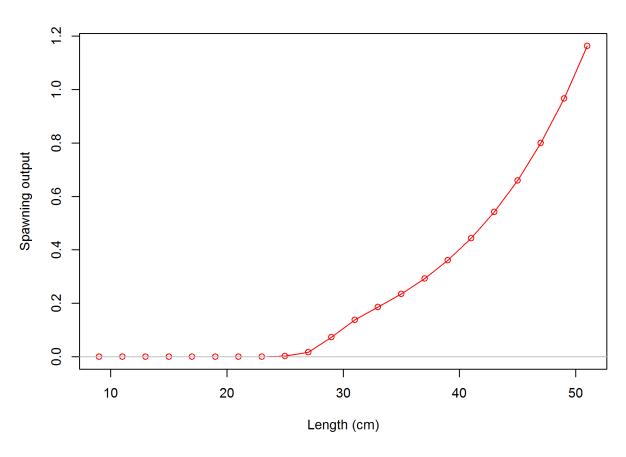
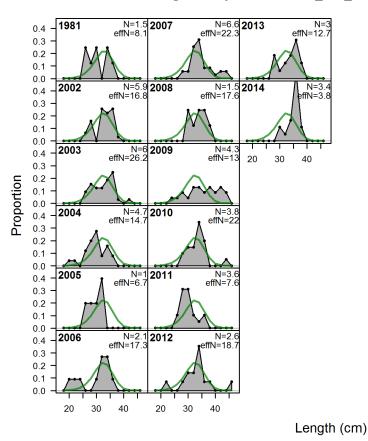


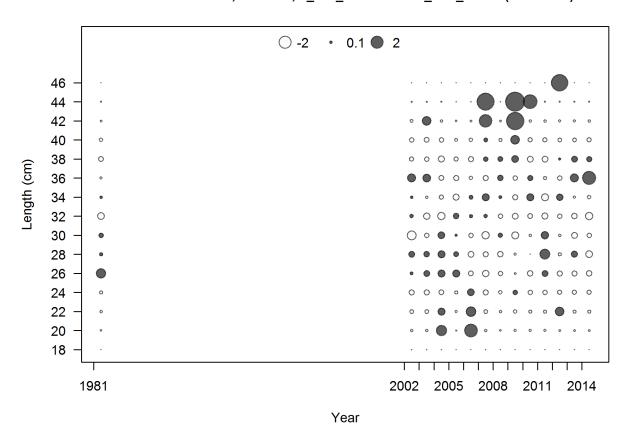
Figure 10: Spawning output at length | fig:mod1_9_bio9_spawningoutput

length comps, retained, 1_WA_SouthernWA_Rec_PCPR

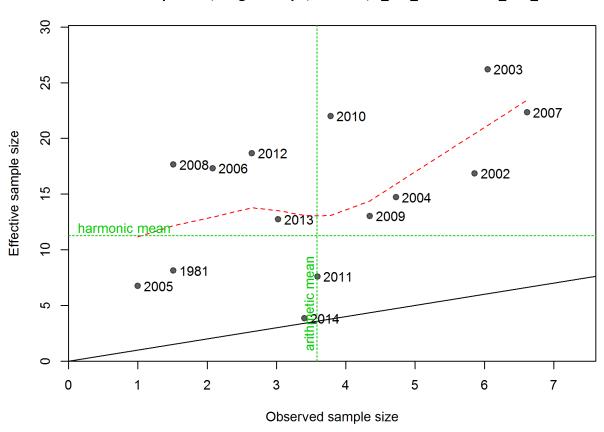


 $\label{eq:figure 11:length comps} Figure \ 11: \ length \ comps, \ retained, \ 1_WA_SouthernWA_Rec_PCPR \ {\tt fig:mod1_1_comp_lenfit_flt1mkt2}$

Pearson residuals, retained, 1_WA_SouthernWA_Rec_PCPR (max=4.76)



N-EffN comparison, length comps, retained, 1_WA_SouthernWA_Rec_PCPR



 $Figure~13:~N_EffN~comparison,~length~comps,~retained,~1_WA_SouthernWA_Rec_PCPR~\\ \\ [fig:mod1_3_comp_lenfit] \\ [fig:mod1_3_comp$

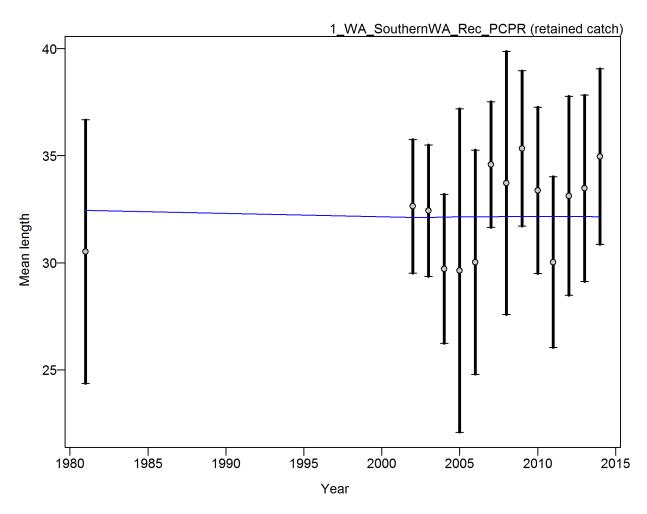
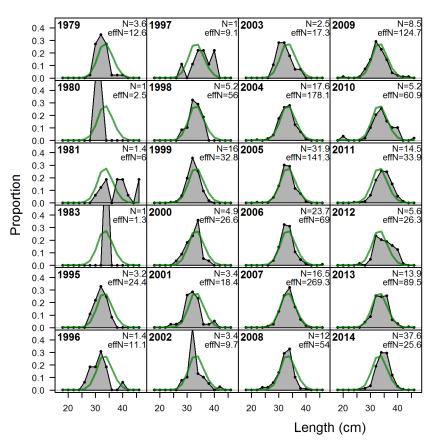


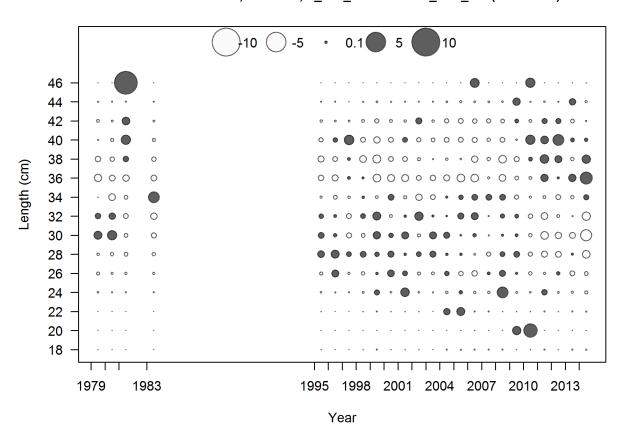
Figure 14: Francis data weighting method TA1.8 1_WA_SouthernWA_Rec_PCPR Suggested sample size adjustment (with 95% interval) for len data from 1_WA_SouthernWA_Rec_PCPR: 0.9991 (0.6704_2.4014) fig:mod1_4_comp_lenfit_data_weighting_TA1.8_1_WA_SouthernWA_Rec_PCPR

length comps, retained, 3_WA_NorthernWA_Rec_PR

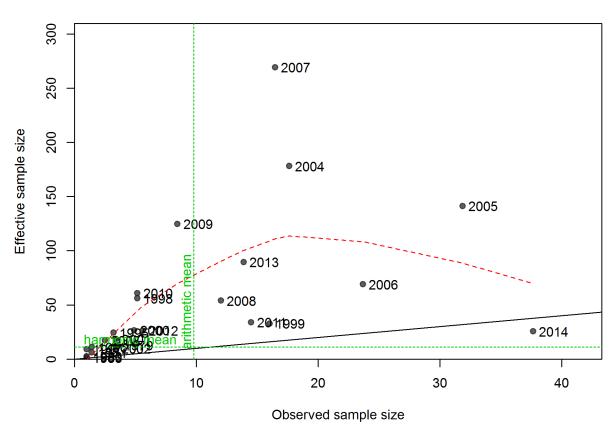


 $Figure~15:~length~comps,~retained,~3_WA_NorthernWA_Rec_PR \\ ~ fig:mod1_5_comp_lenfit_flt3mkt2 \\$

Pearson residuals, retained, 3_WA_NorthernWA_Rec_PR (max=6.82)



N-EffN comparison, length comps, retained, 3_WA_NorthernWA_Rec_PR



 $Figure~17:~N_EffN~comparison,~length~comps,~retained,~3_WA_NorthernWA_Rec_PR~ \\ fig:mod1_7_comp_lenfit_started,~2_WA_NorthernWA_Rec_PR~ \\ fig:mod1_7_comp_lenfit_started,~2_WA_NORTHernWA_R$

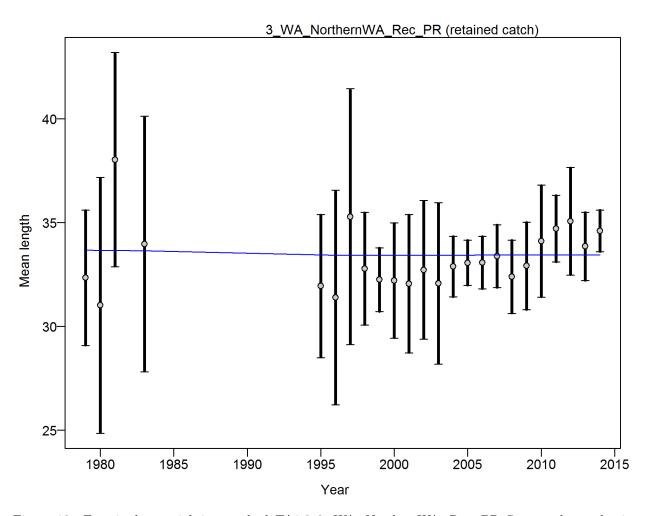
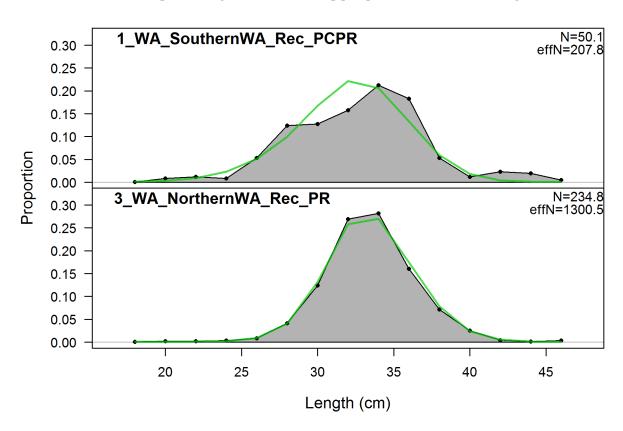


Figure 18: Francis data weighting method TA1.8 3_WA_NorthernWA_Rec_PR Suggested sample size adjustment (with 95% interval) for len data from 3_WA_NorthernWA_Rec_PR: $0.9797 \ (0.6466_2.3089)$ fig:mod1_8_comp_fig:mod1_8_co

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

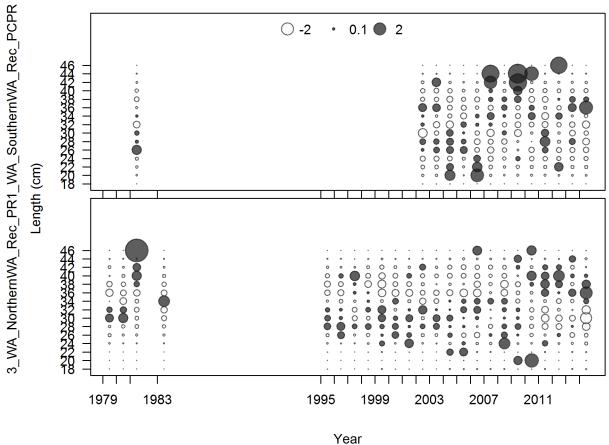


Figure 20: Note: this plot doesn't seem to be working right for some models. Pearson residuals, sexes combined, retained, comparing across fleets Closed bubbles are positive residuals (observed > expected) and open bubbles are negative residuals (observed < expected). fig:mod1_10_comp_lenfit_sex1mkt2_multi-fleet_comparison

Data by type and year

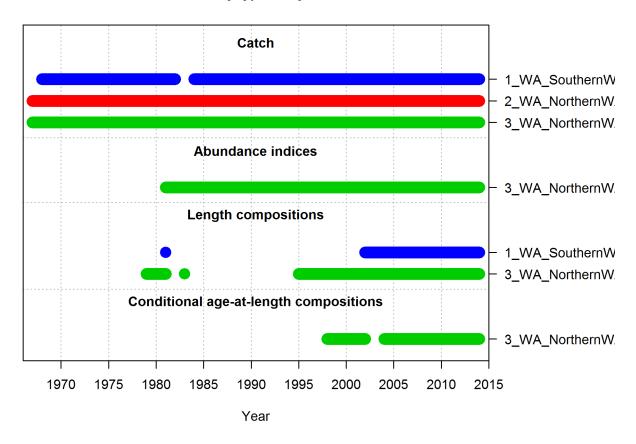


Figure 21: 1-Summary of data sources used in the base case assessment. fig:data_plot

Data by type and year

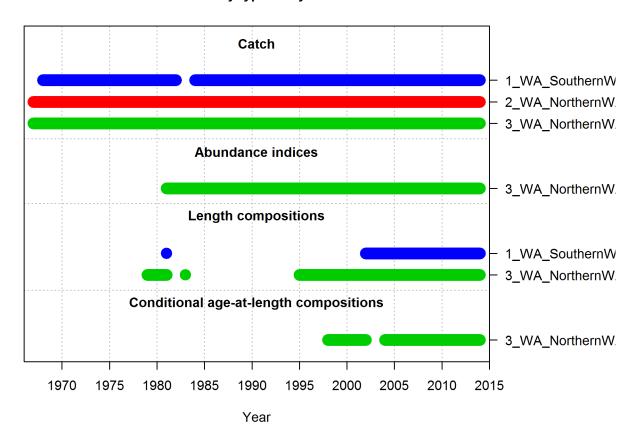


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Data by type and year

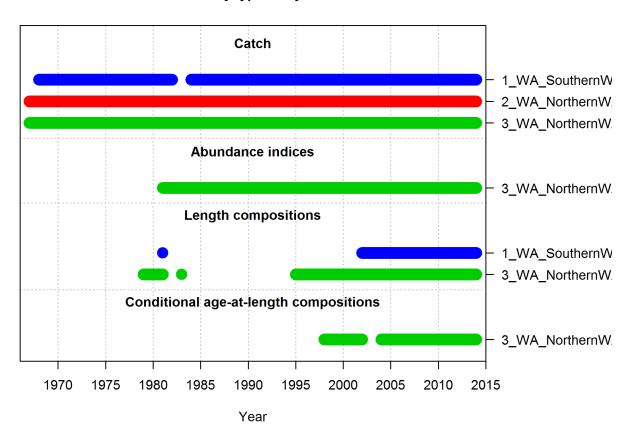


Figure 22: 3-Summary of data sources used in the base case assessment. fig:data_plot