TIM Diagnostics

MODEL STRUCTURE

Match/Mismatch: Matching

Population Structure: Spatial Heterogeneity

Diagnostic Run: NO. Uses OBS values as data inputs

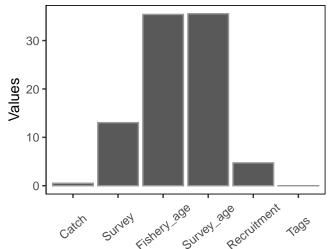
Residual plots represent RELATIVE difference; ((TRUE-ESTIMATED)/TRUE)*100

Run Time of Estimation: 0.0555 minutes

OM Error Parameters

| | Parameter | 1 | 2 |
|---|------------------|------|------|
| 1 | Sigma_Rec | 0.4 | NA |
| 2 | Sigma_Rec_Apport | 0.2 | NA |
| 3 | Sigma_F | 0.2 | 0.2 |
| 4 | Rec_Index_sigma | 0.5 | 0.5 |
| 5 | Survey_Sigma | 0.2 | 0.2 |
| 6 | Catch_Sigma | 0.05 | 0.05 |
| 7 | SIM_N_Catch | 100 | 100 |
| 8 | SIM_N_Survey | 100 | 100 |

Likelihood Components



Model Converged

num gradient component = 8.18356955269281€

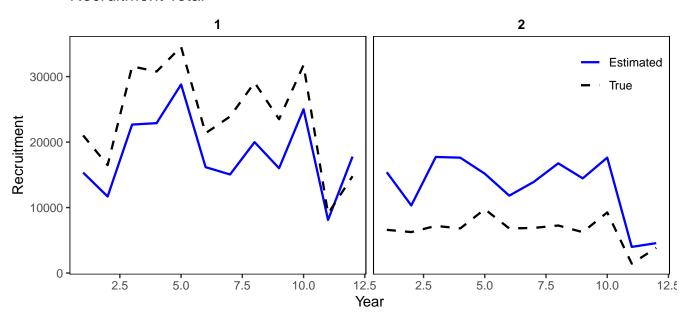
Estimated Parameter Values

| | Parameter | 1 | 2 |
|----|----------------|----------|----------|
| 1 | q | 0.00 | 0.00 |
| 2 | R_ave | 32300.80 | 32300.80 |
| 3 | beta 1 fishery | 5.33 | 5.00 |
| 4 | beta 2 fishery | 1.63 | 1.93 |
| 5 | beta 3 fishery | NA | NA |
| 6 | beta 4 fishery | NA | NA |
| 7 | beta 1 survey | 4.72 | 4.72 |
| 8 | beta 2 survey | 1.65 | 1.65 |
| 9 | beta 3 survey | NA | NA |
| 10 | beta 4 survey | NA | NA |

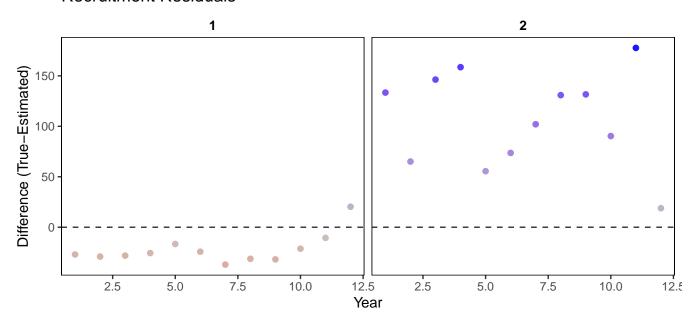
True Parameter Values

| | Parameter | 1 | 2 |
|----|----------------|----------|----------|
| 1 | q | 0.00 | 0.00 |
| 2 | R_ave | 27615.00 | 27615.00 |
| 3 | beta 1 fishery | 1.31 | 2.45 |
| 4 | beta 2 fishery | 3.76 | 3.16 |
| 5 | beta 3 fishery | NA | NA |
| 6 | beta 4 fishery | NA | NA |
| 7 | beta 1 survey | 2.51 | 1.81 |
| 8 | beta 2 survey | 2.31 | 2.64 |
| 9 | beta 3 survey | NA | NA |
| 10 | beta 4 survey | NA | NA |
| | | | |

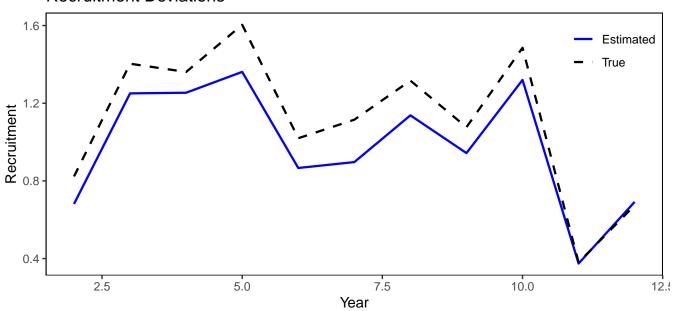
Recruitment Total



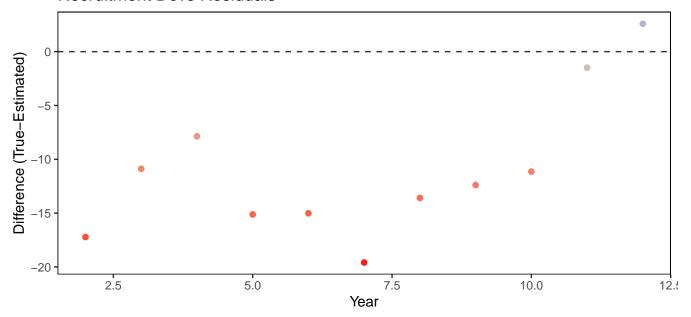
Recruitment Residuals



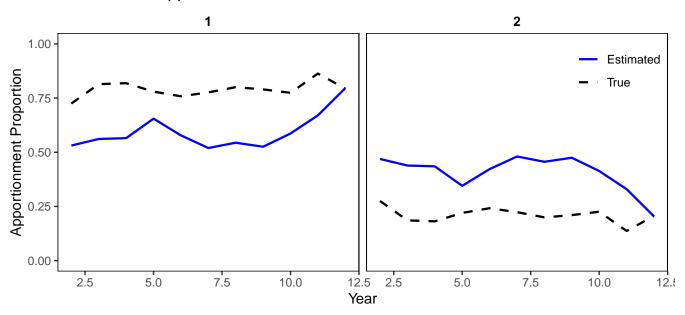




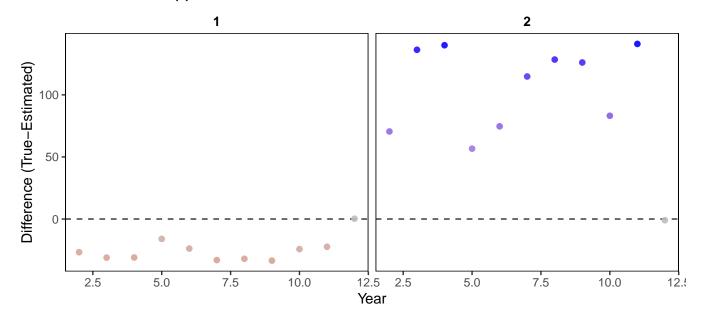
Recruitment Devs Residuals

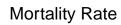


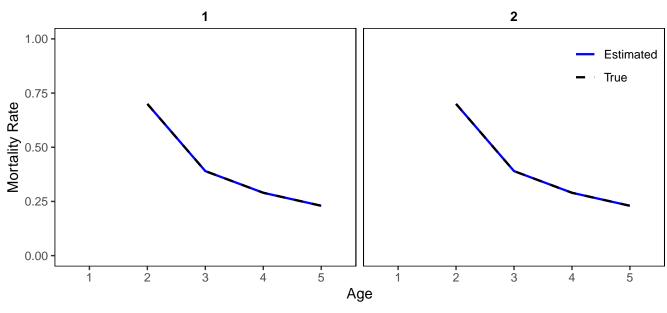
Recruitment Apportionment



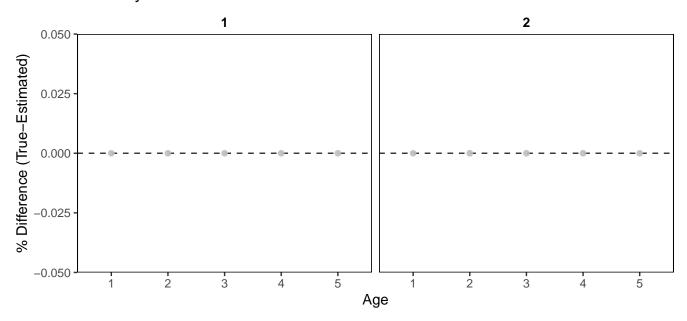
Recruitment Apportionment Residuals



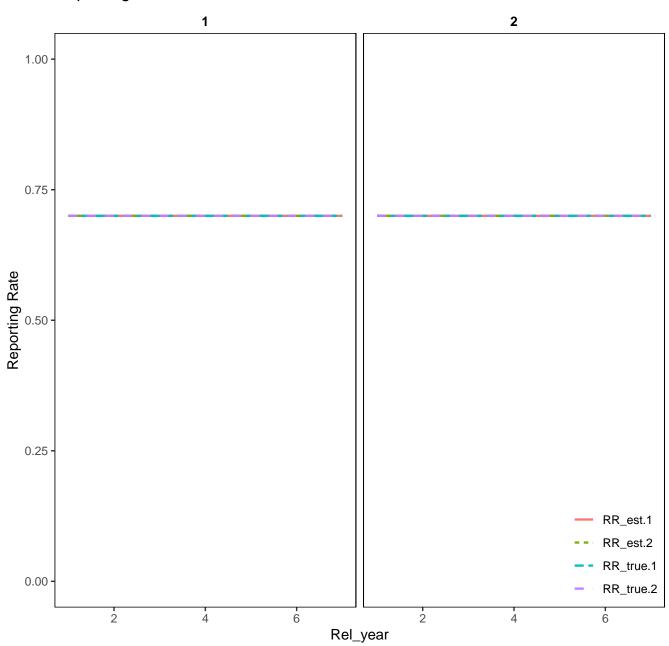




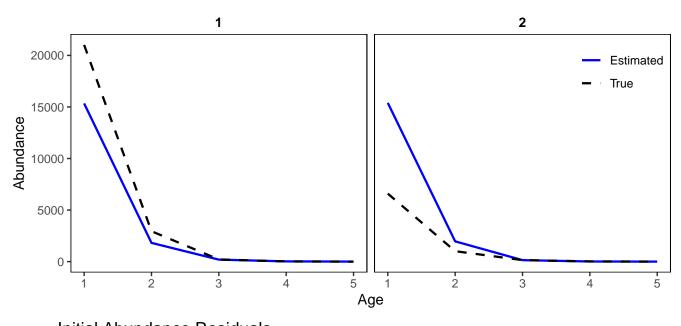
Mortality Rate Residuals



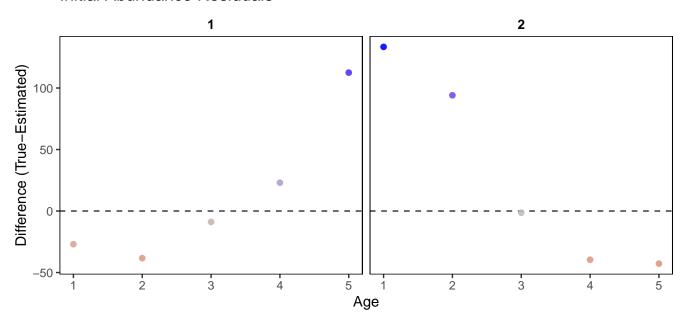




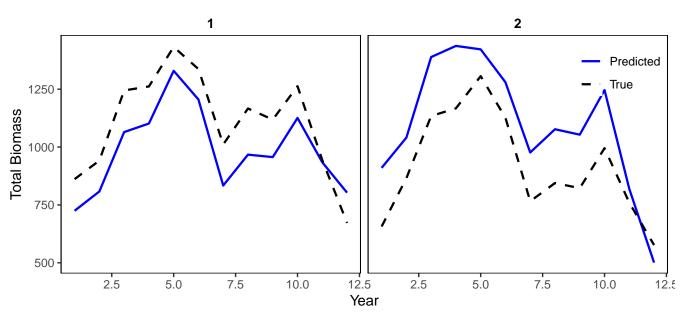
Initial Abundance



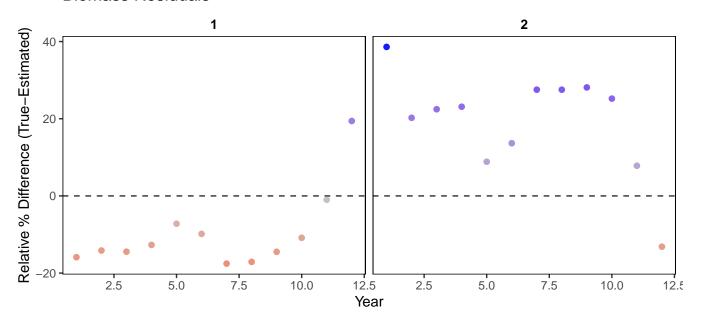
Initial Abundance Residuals

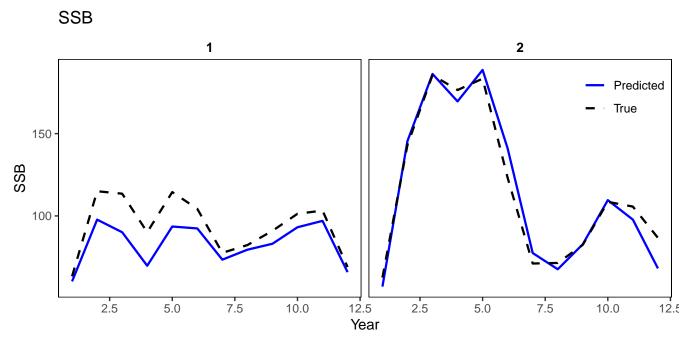


Total Biomass

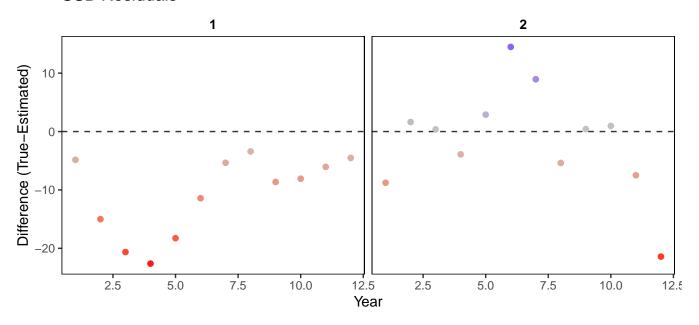


Biomass Residuals

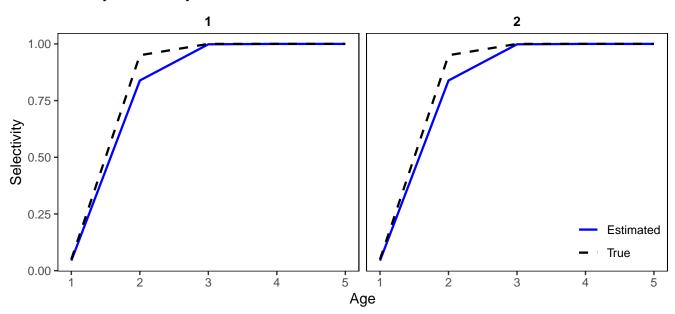




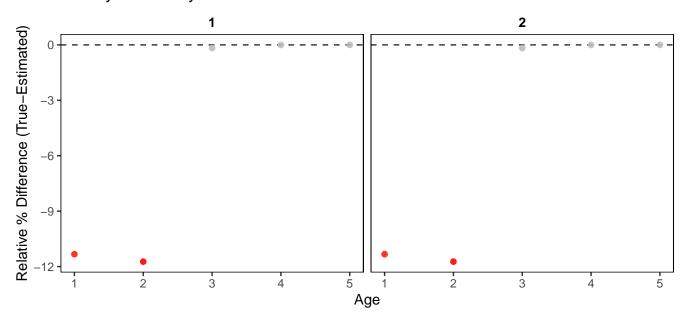


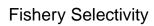


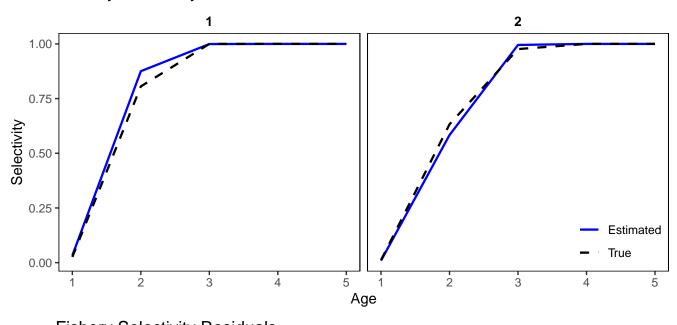




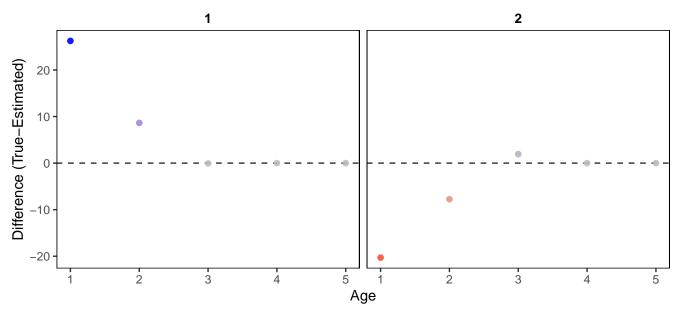
Survey Selectivity Residuals





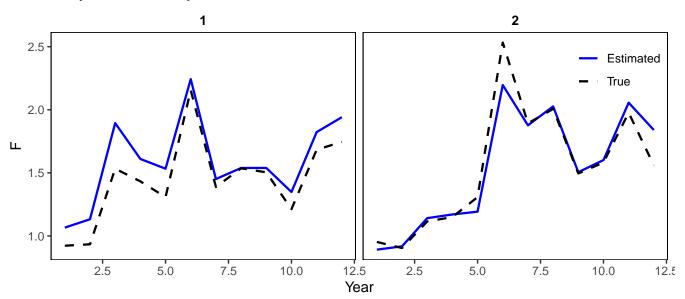


Fishery Selectivity Residuals

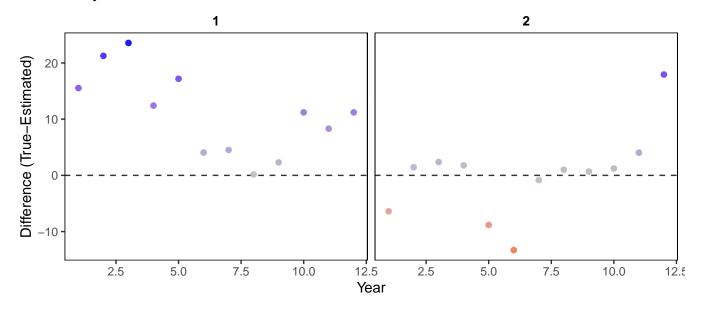


Fishing Mortality

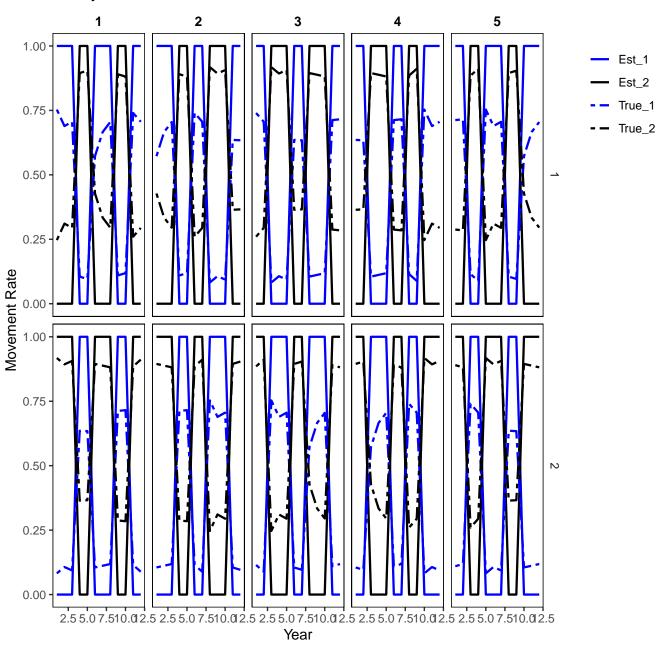




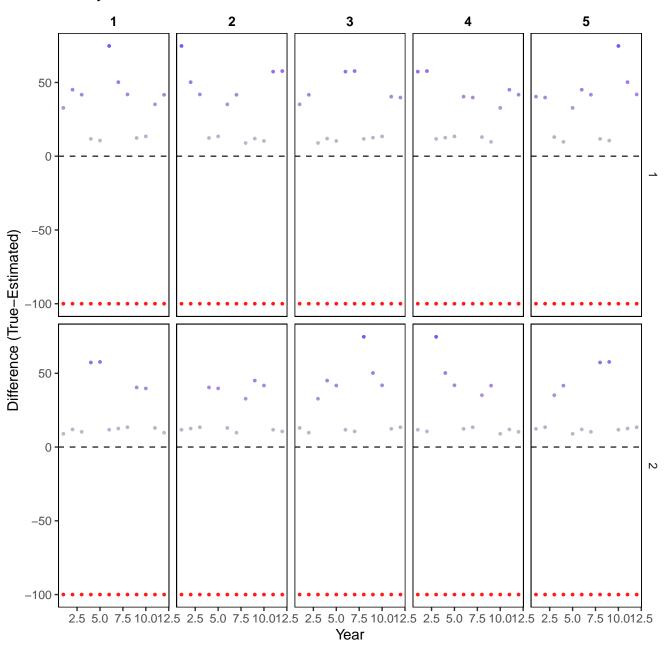
Fully Selected F Residuals

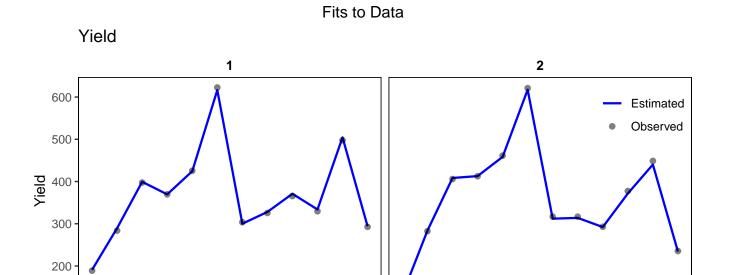






Yearly Movement Rate Residuals





12.5 Year 2.5

5.0

7.5

10.0

12.5

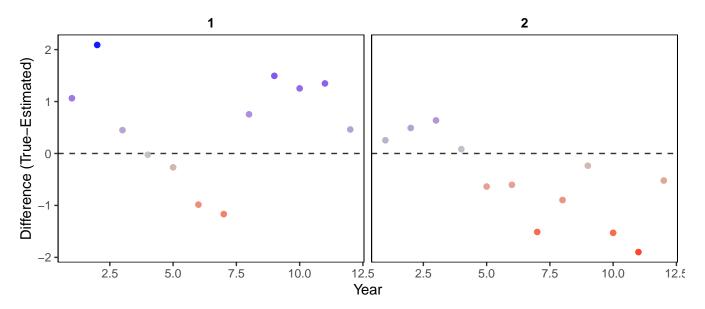


2.5

5.0

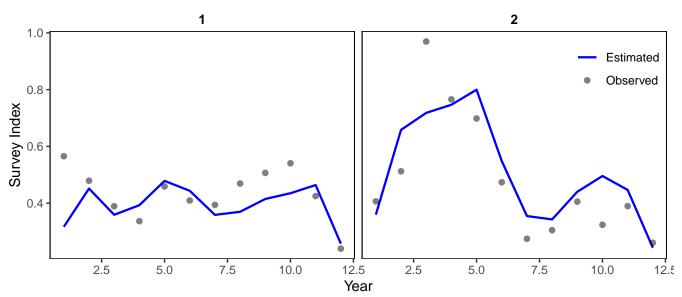
7.5

10.0

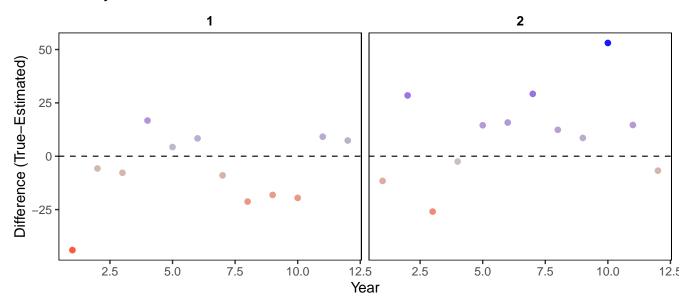


Fits to Data continued





Survey Index Residuals



Fits to Data continued

